

STIC Search Report Biotech-Chem Library

STIC Database Tracking Number: 101115

TO: Ruixiang Li

Location: CM1/10D19/10E18

Art Unit: 1646

Wednesday, August 13, 2003

Case Serial Number: 09/727739

From: Edward Hart

Location: Biotech-Chem Library

CM1-6B02

Phone: 305-9203

edward.hart@uspto.gov

Search Notes

Examiner Li,

Here are the results of the search you requested.

Please feel free to contact me if you have any questions.

Edward Hart



Title: Perfect score: Sequence: Scoring table:

Searched:

OM protein

Rùn on:

Database :

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Human somatostatin relat Somatostatin relat Somatostatin relat Somatostatin relat Somatostatin-28 ($ Rat preprosomatost Human Somatostatin MAPPs p.WMPmp20-(His Human DiTHP polype Swine duodenum elic Somatostatin relat Somatostatin relat Somatostatin-28 pe Somatostatin-28 an Somatostatin-38 pe Somatostatin-38 pe Somatostatin-38 pe Somatostatin-38 pe Somatostatin-38 pe Somatostatin-38 pe Somatostatin analo Somatostatin analo Somatostatin analo Somatostatin analo Radioactively tagg Somatostatin-38 an Somatostatin-38 pe Somatostatin-38 pe Somatostatin-38 pe Somatostatin-38 pe Ralinbow trout soma
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Somatostatin-28 pe
Somatostatin-28 an
Somatostatin antig
Somatostatin antig
                                                                     Peptide hormone so
Somatostatin relat
                                                                                                                     Mammalian somatost
Somatostatin analo
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/note= "ppss-II'' pre-sequence"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             26.111
/note= "Mature PPSS-II''"
87.97
/note= "PPSS-II'' pro-sequence"
88.111
/note= "Prosomatostatin II'"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1. 25
/note= "Signal peptide"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAU07668 standard; Protein; 111 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         04-DEC-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Oncorhynchus mykiss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAU07668;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Peptide
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Sequence of prepro
Rainbow trout prep
Somatostatin-28 an
Analogue of angler
Rainbow trout prep
Sequence of prepro
Sequence of prepro
Sequence of somato
Somatostatin relat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                    August 13, 2003, 14:45:37; Search time 46.7949 Seconds (without alignments) 84.799 Million cell updates/sec
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Description
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A_Geneseq_19Jun03:*

| A_SID51/gcdata/geneseq/geneseqp-emb1/AA1980.DAT:*
| SID51/gcdata/geneseq/geneseqp-emb1/AA1981.DAT:*
| SID51/gcdata/geneseq/geneseqp-emb1/AA1981.DAT:*
| SID51/gcdata/geneseqf/geneseqp-emb1/AA1982.DAT:*
| SID51/gcdata/geneseqf/geneseqp-emb1/AA1982.DAT:*
| SID51/gcdata/geneseqf/geneseqp-emb1/AA1982.DAT:*
| SID51/gcdata/geneseqf/geneseqp-emb1/AA1982.DAT:*
| SID51/gcdata/geneseqf/geneseqp-emb1/AA1987.DAT:*
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/SIDSI/gcddata/geneseq/geneseqp-embl/AA1996.DAT:
/SIDSI/gcddata/geneseq-geneseqp-embl/AA1996.DAT:
/SIDSI/gcddata/geneseq-geneseqp-embl/AA1996.DAT:
/SIDSI/gcddata/geneseq-geneseqp-embl/AA1999.DAT:
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/SIDSI/gcddata/geneseq-geneseqp-embl/AA2000.DAT:
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SIDSI/gcgdata/geneseg/genesegp-embl/AA199
SIDSI/gcgdata/geneseg/genesegp-embl/AA199
SIDSI/gcgdata/geneseg/genesegp-embl/AA199
                                          GenCore version 5.1.6
Copyright (c) 1993 - 2003 Compugen Ltd.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                               1107863 segs, 158726573 residues
                                                                                                                                                                                                                                                                                                                                       SVDNLPPRERKAGCKNFYWKGFTSC 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SUMMARIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                22 AAU07668
3 AAP20029
22 AAU07667
7 AAP61714
110 AAP90989
22 AAP20028
3 AAP20028
3 AAP20198
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                                                                                                                             - protein search, using sw model
                                                                                                                                                                                                                                                                                                                                                                                          BLOSUM62
Gapop 10.0 , Gapext 0.5
                                                                                                                                                                                                                                                                                 US-09-727-739B-16
147
1 SVDNLPPRERKAGCKN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Minimum DB seq length: 0
Maximum DB seq length: 2000000000
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DB

Length

Ouery Match

112..125 /label- Somatostatin II

81EP-0303825 80US-0181046

Location/Qualiflers

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esultant precursor (pro S1) is subsequently cleaved to yield S1 itself. The prepeptide portion of prepro S1 is probably about 20 asses long. Translation of plaS2 predicts the sequence of a 125
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           preprosomatostatin-2, prosomatostatin-2 and somatostatin-2; encoding them. The translation of somatostatin mRNA yields precursor (prepro SI) containing a signal peptide which may released during the translatinto the endoplasmic reticulum.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New somatostatin and precursors - produced by transformed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           eptide which surprisingly contains a 14 AA sequence at erminus which differs from 51 by only 2 AAs, and is ter comatostatin 2 (82)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Updated on 16-AUG-2002 to add missing OS field.) Updated on 25-MAR-2003 to correct PA field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Ouery Match 93.9%; Score 13%; DB 3; Best Local Similarity 92.0%; Pred. No. 5.5e-13; Matches 23; Conservative 1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                           Hobart P, Crawford R, Pictet RL, Rutter WJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The inventors claim preprosomatostatin-1,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 SVDNLPPRERKAGCKNFYWKGFTSC 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example; Fig 3; 50pp; English.
                                                                                                                                                                                                                                                                                                                                                                                       (REGC ) UNIV CALIFORNIA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1982-18113E/10.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 125 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      N-PSDB; AAN20034.
                                                                                                                                                                                                                                                                                                                                       25-AUG-1980;
                                                                                                                                                                                                                                      03-MAR-1982.
                                                                                                                                                                                     EP46669-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAU07667;
                                                                                                             Protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The invention traines of preprosomatostatin (PPSS-I) and/or a portion of preprosomatostatin (PPSS-I) and/or a portion of preprosomatostatin (PPSS-II) and/or a portion of preprosomatostatin II (PPSS-II). The protein sequences and their associated polynucleotides are useful for identifying modified somatostatin polypeptide which functions as a somatostatin agonist useful for research, therapeutics or diagnostics, including medical and veterinary applications. The wild-type somatostatin and its modified version are useful for treating hypersecretion from endocrine tumours in the pituitary (e.g. acromegaly) or gastroenteropancreatic tissues (e.g. gastrouna, glucagnoma, carcinold syndrome), to cause tumour shrinkage through their effects on cell proliferation and apoptosis and as adjuncts in the treatment of diabetes mellitus via inhibition of growth hormone
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     invention relates to an Oncorhynchus mykiss somatostatin polypeptide taining a portion of preprosomatostatin I (PPSS-I) and/or a portion of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    their secretory products
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  with acquired immunodeficiency syndrome (AIDS) and various all disorders (e.g. epilepsy, Alzheimer's disease and 's disease) and somatostatin antagonists are effective in the 5s such conditions. Nucleic acids encoding the polypeptides argene therapy and fusion peptides can be targeted to neoplasms
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   equence represents O. Mykiss PPSS-II'' protein.
The features for this sequence are specifically claimed in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New somatostatin polypeptides derived from Oncorhynchus mykiss, u
for-treating diabetes mellitus, acromegaly, gassrinoma, acquired
immunodeficiency syndrome and neurological disorders -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                somatostatin secretion
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        / Match 100.0%; Score 147; DB 22; Length 111; Local Similarity 100.0%; Pred. No. 2.1e-14; nes 25; Conservative 0; Mismatches 0; Indels 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Somatostatin; growth hormone; peptide hormone; secretion
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence of preprosomatostatin-2 encoded on plaS2.
                           "Dibasic cleavage site"
96..97
/note- "Dibasic cleavage site"
98.111
/note- "SS-14 variant peptide"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 the treatment of diabetes mellitus via glucagon. In addition, dysfunctional
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   87 SYDNLPPRERKAGCKNFYWKGFTSC 111
                                                                                                                                                                                                                                                                                                                                                                                       Kittelson JD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 SVDNLPPRERKAGCKNFTWKGFTSC 25
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   therapy and fusion
tases, inhibiting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 1; Fig 3; 52pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 treatment of such conditions
                                                                                                                                                                                                                             01-DEC-2000; 2000CA-2325169.
                                                                                                                                                                                                                                                                                   99US-0168934
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (updated)
(updated)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                       Sheridan MA, Moore CA,
                                                                                                                                                                                                                                                                                                                                    (NDSU-) NDSU RES FOUND
                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2001-425997/46.
N-PSDB; AAS12935.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 111 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            his sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       specification
                                                                                                                               CA2325169-A1
                                                                                                                                                                                                                                                                                   03-DEC-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  25-MAR-2003
16-AUG-2002
14-AUG-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                neurological
                                                                                                                                                                               03-JUN-2001.
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Best Local
                                                   Peptide
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and DNA

its carboxy

Length 125;

1; Indels

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Rainbow trout; somatostatin; preprosomatostatin; hypersecretion; PPSS-I; PPSS-II; PPSS-II; pepS-II; pedocrine tumour; pitulitary gland; glucagonoma; AIDS; gastroenteropanoreatic tissue; acromegaly; gastrinoma; diabetes mellitus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Rainbow trout preprosomatostatin II (PPSS-II') polypeptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           carcinoid syndrome; cell proliferation; apoptosis; growth h
glucagon; acquired immunodeficiency syndrome; neurological
epilepsy; Alzhelmer's disease; Huntington's disease; neurop
neoplasm; metastasis; gene therapy; antidiabetic; noofropio
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            osteopathic;
101 STNNLPPRERKAGCKNFTWKGFTSC 125
                                                                                                                                                                                                                                           AAU07667 standard; Protein; 115 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       anti-human immunodeficiency virus;
                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               neoplasm; metastasis;
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ZKEKEZEZ ZKEKEEZ

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Somatostatin-28; insulin-selective; insulinoma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (SALK ) SALK INST. BIOLOGICAL STUDIES.
                                                        AAP61714 standard; Protein; 28 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 31-AUG-1984; 84US-0646610.
01-APR-1987; 87US-0033295.
                                                                                                                                                                                                                                   Somatostatin-28 analogue.
                                                                                                                                                                                                                                                                                                                                                                                                                    Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 16-AUG-1985;
                                                                                                                                                        25-MAR-2003
28-JUL-1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              05-MAR-1986.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           EP173527-A.
                                                                                                        AAP61714;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The invention relates to an Oncorhynchus mykiss somatostatin polypeptide containing a portion of preprosomatostatin I (PPSS-I) and/or a portion of preprosomatostatin II (PPSS-II). The protein sequences and their associated polynucleotides are useful for identifying modified somatostatin polypeptide which functions as a somatostatin agonist useful for research, therapeutics or diagnostics, including medical and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              sequified immunodeficiency syndrome (AIDS) and various isorders (e.g. epilepsy, Alzheimer's disease and sisease) and somatostatin antagonists are effective in the uch conditions. Nucleic eacids encoding the polypeptides are therapy and fusion peptides can be targeted to neoplasms istases, inhibiting the release of their conditions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            uitary (e.g. acromegaly) or gastroenteropancreatic tissues (e.g. ma, Julucagonoma, carcinoda syndrome), to cause tumour shrinkage their effects on cell proliferation and apoptosis and as adjuncts treatment of diabetes mellitus via inhibition of growth hormone
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               terinary applications. The wild-type somatostatin and its modified reson are useful for treating hypersecretion from endocrine tumours in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ence represents O. MyKiss PPSS-II' protein. features for this sequence are specifically claimed in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New somatostatin polypeptides derived from Oncorhynchus mykiss, u
for treating diabetes mellitus, acromegaly, gastrinoma, acquired
immunodeficiency syndrome and neurological disorders -
                                                                                                                                                                                                                                                                                                                                100..101
/note- "Dibasic cleavage site"
                                                                                                                                                                                                                    /mote- "PPSS-II' pro-sequence"
                                                                                                                                                                                                                                                                                                                                                                                                           /note= "SS-14 variant peptide"
                                                                                            "PPSS-II' pre-sequence
                                                                                                                                                                                                                                                                       /more- "Prosomatostatin II'"
                                                                                                                          76..115
/note= "Mature PPSS-II'"
                          ...25
'note= "Signal peptide"
                                                                                                                                                                                                     note- "Encoded by CAA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sheridan MA, Moore CA, Kittelson JD;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 2; Fig 3; 52pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          01-DEC-2000; 2000CA-2325169.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                99US-0168934
                                                                                                                                                                                                                                                                                                                                                                                      ..115
                                                                             ...87
'note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (NDSU-) NDSU RES FOUND.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2001-425997/46.
N-PSDB; AAS12934.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ul in gene theragitheir interestable
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        115 AA;
                                                                                                                                                                          Misc-difference
                                                                                                                                                                                                                                                                                                                                Cleavage-site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   03-DEC-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     specification
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             03-JUN-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence
Key
Peptide
                                                                                                                                                                                                                             Pept1de
                                                                                                                                                                                                                                                                                                                                                                                      Peptide
                                                                       Protein
                                                                                                                          Protein
                                                                                                                                                                                                                                                                                Peptide
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Location/Qualifiers

/label- Hyl, Lys

85EP-0305867

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Angler fish somatostatin-28 and analogue and fragment - useful in
inhibiting insulin secretion in insulinoma.
                                                                                                                                                                                                                                                                                                                                                                                                              Score 132; DB 7; Length 28;
Pred. No. 9.4e-13;
1; Mismatches 2; Indels
                                                                                                                                                                                                   The protein sequence is an insulin-selective analogue of anglerfish somatostatin-28, which is more potent than somatostatin-14 or somatostain-28 in inhibiting insulin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Somatostatin 28; SS-28; analogue; insulin secre inhibitor; insulinoma; gastric acid secretion;
                                                                                                                                                                                                                                                                           secretion for treatment of insulinoma. (Updated on 25-MAR-2003 to correct PR field.) (Updated on 25-MAR-2003 to correct PA field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Analogue of anglerfish somatostatin 28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 SVDNLPPRERKAGCKNFYWKGFTSC 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAP90989 standard; peptide; 28 AA.
                                                                                                                                                          Claim 4; Page 18; 19pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 89.8%;
Best Local Similarity 88.0%;
Matches 22; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (updated)
(first entry)
                                          WPI; 1986-063363/10.
Spiess J, Noe BD;
                                                                                                                                                                                                                                                                                                                                                                      28 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         25-MAR-2003
08-JUN-1990
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Anglerfish
                                                                                                                                                                                                                                                                                                                                                                      Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAP90989;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 5
AAP90989
ID AAP9
XX
AC AAP9
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Match 91.2%; Score 134; DB 22; Length 115; Local Similarity 95.7%; Pred. No. 2e-12; conservative 1; Mismatches 0; Indels (

Query Match

Matches

å . ପ୍ର

3 DNLPPRERKAGCKNFYWKGFTSC 25

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01-DEC-2000; 2000CA-2325169.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sheridan MA, Moore CA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (NDSU-) NDSU RES FOUND
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2001-425997/46...
N-PSDB; AAS12933.
                                                                                                                                                                                                                                  Cleavage-site
                                                                                                                                                                                                                                                                                                                                              CA2325169-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             03-DEC-1999;
                                                                                                                                                                                                                                                                                                                                                                                      03-JUN-2001.
                                                                                                                                                                                                                                                                              Peptide
                                                             Protein
                                                                                                        Protein
                                                                                                                                              Peptide
                                                                                                                                                                                          Peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                It is called aSS-28 because it is an analogue of anglerfish somatostatin (SS-28). It is more potent than either somatostatin id (SS-14) or SS-28 at inhibiting insulin secretion for the treatment of insulinoma. The for inhibiting insulin secretion for the pancreas. aSS-28 and asS-18 is useful for inhibiting insulin secretion by the pancreas. aSS-28 and aSS-28 may be useful for decreasing gastric acid secretion and influencing them or Their reduced linear forms, wherein the disulphide (Updated on 25-MAR-2003 to correct by field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Angler fish somatostatin-28 and fragments -
useful in inhibiting insulin secretion and insulinoma
                                                     /note-"Bonded to Cys-28"
                                                                                              'note-"Bonded to Cys-17"
                                                                                                                                                                             /note="Also claimed"
                                                                                                                                                                                                                                                                                                                                                                                                                               (SALK ) SALK INST BIOLOGICAL STUDIES.
Location/Qualifiers
17
                                                                                                                                            label-Lys, Hyl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 1; page 65; 8pp; English.
                                                                                                                                                                                                                                                                                                                                                             87US-0033295.
                                                                                                                                                                                                                                                                                                                    87US-0033295
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 1989-113910/15.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Spiess J, Noe BD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        28 AA;
                                                                                                                     Misc-difference
               Key
Disulfide-bond
                                                                         Disulfide-bond
                                                                                                                                                                                                                                                                                                                                                             01-APR-1987;
31-AUG-1984;
                                                                                                                                                                                                                                                                                                                    01-APR-1987;
                                                                                                                                                                                                                             US4816438-A.
                                                                                                                                                                                                                                                                         28-MAR-1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence
                                                                                                                                                                Region
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.100 te= "Dibasic cleavage site"

/note-

101..114 /note= "SS-14 peptide"

Kittelson JD;

99US-0168934

"ppss-I pro-sequence" "Prosomatostatin I"

/note-

89.114 /note= "P 99.100

'note- "PPSS-I pre-sequence"

/ucc. 25..114 /~rte= "Mature PPSS-I"

'note- "Signal peptide"

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somatostatin polypeptide which functions as a somatostatin agonist useful for research, therapeutics or diagnostics, including medical and veterinary applications. The wild-type somatostatin and its modified version are useful for treating hypersecretion from endocrine tumours in the pituitary (e.g. acromegaly) or gastroenteropancreatic tissues (e.g. gastrinoma, glucagnoma, carcinoid syndrome), to cause tumour shrinkage through their effects on cell proliferation and apoptosis and as adjuncts in the treatment of diabetes mellitus via inhibition of growth hormone
                                                                                                                                                                                                              The invention relates to an Oncorhynchus mykiss somatostatin polypeptide containing a portion of preprosomatostatin I (PPSS-I) and/or a portion of preprosomatostatin II (PPSS-II). The protein sequences and their associated polynucleotides are useful for identifying modified somatostatin polypeptide which functions as a somatostatin agonist useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     of their secretory products.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              are effective in the
New somatostatin polypeptides derived from Oncorhynchus mykiss, useful for treating diabetes mellitus, acromegaly, gastrinoma, acquired immunodeficiency syndrome and neurological disorders -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   omatostatin secretion is syndrome (AIDS) and various
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               . Nucleic acids encoding the polypeptides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  nce represents O. Mykiss PPSS-I protein.
features for this sequence are specifically claimed in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ;
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 114;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           somatostatin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         neurological disorders (e.g. epilepsy, Alzheimer's
Huntington's disease) and somatostatin antagonists
treatment of such conditions. Nucleic acids encodin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 107; DB 22;
Pred. No. 2.4e-08;
1; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           In addition, dysfunctional :
h acquired immunodeficiency
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAP20028 standard; Protein; 121 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LAPRERKAGCKNFFWKTFTSC 114
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             5 LPPRERKAGCKNFYWKGFTSC 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 72.8%; Soc
Best Local Similarity 85.7%; Pro
Matches 18; Conservative 1;
                                                                                                                                            Claim 1; Fig 2; 52pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      This sequence represents
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 114 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        specification
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 7
AAP20028
ID AAP2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ö
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Rainbow trout; somatostatin; preprosomatostatin; hypersecretion; PPSS-1; PPSS-11; PPSS-11; endocrine tumour; pitulitary gland; glucagonoma; AIDS; gastroenteropanoreatic tissue; acromegaly; gastrinoma; diabetes mellitus; carcinoid syndrome; cell proliferation; apoptosis; growth, hormone; apuncagon; acquired immunodeficiency syndrome; neurological disorder; HIV; epilepsy; Alzheimer's disease; Huntington's disease; neuroporotective;

Rainbow trout preprosomatostatin I (PPSS-I) polypeptide.

04-DEC-2001 (first entry)

AAU07666 standard; Protein; 114 AA

PESULT 6

Query.Match

89.8%; Score 132; DB 10; Length 28;
Best Local Similarity 88.0%; Pred. No. 9.4e-13;
Matches 22; Conservative 1; Mismatches 2; Indels

å 셤 neoplasm; metastasis; gene therapy; antidiabetic; nootropic; canti-human immunodeficiency virus; osteopathic; anticonvulsant

Location/Qualiflers

Oncorhynchus mykiss.

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us-09-727-739b-16.rag

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Protection, endogenous therapeutic peptide; peptidase; conjugation; blood component; modification; succinimidyl; maleimido group; amino; hydroxyl; thiol; hormone; growth factor; neurotransmitter.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  inhibiting release of growth hormone; they also inhibit basal and stimulated insulin and glucagon sectretion. (D-Trp(22))-SS-28 exhibits very substantial increases in potency w.r.t. inhibition of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       5 or (D-Trp(22))-SS-28 and derivs, and SS-25 derivs, then somatostatin in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The inventors claim a pharmaceutical compsn. which comprises (1) synthetic somatostatin. 18 (SS-28), SS-25 or (D-Trp(22)), SS-28 and (11) a liq. or solid carier, and SS-28 derive, and SS-28 derives and the compsn. and derive, are more potent then somatostatin in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Extended somatostatin analogues - with increased inhibition of growth hormone release
Somatostatin; hormone; growth hormone release; inhibition.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 105; DB 3; Length 25;
Pred. No. 1e-08;
2; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       growth hormone secretion. (Updated on 16-AUG-2002 to add missing OS field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Bohlen P, Brazeau
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Somatostatin related peptide SEQ ID NO:191.
                                                                                                             Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAB91017 standard; Peptide; 25 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             5 LPPRERKAGCKNFYWKGFTSC 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     5 MAPRERKAGCKNFFWKTFTSC 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 6; Column 12; 7pp; English.
                                                                                                                                                                                                                                                                                                                                                                 80US-0159801.
                                                                                                                                                                                                                              /label- C-OH
                                                                                                                                                                                                                                                                                                                                                                                                              80US-0159801.
                                                                                                                                                     _label= H-S
14..25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 71.4%;
Best Local Similarity 81.0%;
Matches 17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Guillemin RCL, Esch FS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                            (SALK-) SALK INST BIOLOG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 1982-19801E/10.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          25 AA;
                                                                                                                                                                           Disulfide-bond
Modified-site
                                                                                                           Key
Modified-site
                                                                                                                                                                                                                                                                                                                                                                                                              14-JUN-1980;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO200069900-A2
                                                                                                                                                                                                                                                                                                                                                                 14-JUN-1980;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens.
Synthetic.
                                                                                                                                                                                                                                                                      US4316891-A.
                                                                                                                                                                                                                                                                                                                   23-FEB-1982.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              22-JUN-2001
                                        Mammalia.
Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  23-NOV-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAB91017;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   g
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ind the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                esultant precursor (pro SI) is subsequently cleaved to yield SI their Pre-prepetide portion of prepro SI is probably about 20.25 asses long. Translation of plas2 predicts the sequence of a 125 Amely which surprisingly contains a 14 AA sequence at its carboxy erminus which differs from SI by only 2 AAS, and is termed compatostatin 2 (S2).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     and DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ncoding them. The translation of somatostatin mrNa yields recursor (prepro 3) contraining a signal peptide which may eleased dufing the transit into the endoplasmic reticulum.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 121;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New somatostatin and precursors - produced by transformed microorganisms
                                                                                                                                                                                   Somatostatin; growth hormone; peptide hormone; secretion
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Ouery Match 72.8%; Score 107; DB 3; Length 12
Best Local Similarity 85.7%; Pred. No. 2.6e-08;
Matches 18; Conservative 1; Mismatches 2; Indels
                                                                                                                                        Sequence of preprosomatostatin-1 encoded on plasi.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (Updated on 16-Aug-2002 to add missing OS field.) (Updated on 25-MAR-2003 to correct PA field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Rutter WJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The inventors claim preprosomatostatin-1,
                                                                                                                                                                                                                                                                                Location/Qualifiers
108..121
/label= Somatostatin I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          somatostatin-25 analoque
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Hobart P, Crawford R, Pictet RL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         101 LAPRERRAGCKNFFWKTFTSC 121
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           5 LPPRERKAGCKNEYWKGFTSC 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAP20198 standard; Protein; 25 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example; Fig 3; 50pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    81EP-0303825.
                                                  (updated)
(updated)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 80US-0181046
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(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (REGC ) UNIV CALIFORNIA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               eprosomatostatin-2,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 1982-18113E/10.
N-PSDB; AAN20033.
                                                                                                                                                                                                                                   Lophius americanus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 121 AA;
                                             25-MAR-2003
16-AUG-2002
14-AUG-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  sacoding them
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 21-AUG-1981;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              25-AUG-1980;
                                                                                                                                                                                                                                                                                                                                                                                                                   03-MAR-1982
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 16-AUG-2002
14-AUG-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence of
                                                                                                                                                                                                                                                                                                                                                                   EP46669-A.
     AAP20028;
                                                                                                                                                                                                                                                                                   Key
Protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAP20198;
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Gaps

us-09-727-739b-16.rag

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the risk factors of syndrome X of Reaven (hyperinsulinaemia syndrome). It comprises somatostatin, diazoxide, cyclothazide (or an analogue of one of these) or metformin as the active ingredient. The composition reduces resistance to insulin, and so treats and prevents all the associated risk factors at once. The risk factors are hypertension, dyslipidaemia fratsed trigiyceride and LDL levels with reduced HDL levels), shorter
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   disease, impotence due to peripheral vascular disease, hypercoagulation (e.g. renal vein thrombosis), obesity and glucose intolerance. The present sequence represents a specifically claimed somatostatin analogue.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         cerebrovascular disorders, intermittent claudication, ischaemic bowel disease, importence due to peripheral vascular disease, hypercoagulation (e.g. renal vein thrombosts), obesity and glucose intolerance. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         coagulation fine due to increased Plasmainogen Activetor Inhibitor 1 levels, core obesity, glucose intolerance hyperinsulinaemia. The composition reduces the incidence of ischaemic heart disease. The composition reduces the incidence of ischaemic heart disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            neuroendocrine cells, e.g. of
                                                                                                                 Composition for treatment of the risk factors of syndrome X of Reaven - (hyperinsulinaemia syndrome) comprises somatostatin, diazoxide, cyclothiazide (or their analogues) and/or metformin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      71.4%; Score 105; DB 19; Length 28;
81.0%; Pred. No. 1.1e-08;
Live 2; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           thyroid stimulating hormone; octreotide; cell-base glucose-stimulated insulh secretion; SSTRV; somat mouse sometostatin receptor type V gene; diabetes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       5 LPPRERKAGCKNFYWKGFTSC 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        8 MAPRERKAGCKNFFWKTFTSC 28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAY28703 standard; peptide; 28 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New defined medium for culture of insulin-secreting cells
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             somatostatin SS-28 hormone;
                                                                                                                                                                                                                                                  Claim 42; Page 41; 45pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mouse somatostatin SS-28 hormone.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           99WO-US00633
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         98US-0087848
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   98US-0071193
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity 81.0%
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (BETA-) BETAGENE INC.
                                                                  WPI; 1998-271636/24.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Quaade C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               28 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mus musculus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      W09935242-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       12-JAN-1998;
12-JAN-1998;
12-JAN-1998;
03-JUN-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        07-0CT-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         03-JUN-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                15-JUL-1999.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Clark SA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAY28703;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mouse
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 g
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  reactive group (II) (e.g. succinimisty) and malehinido groups) attached to a less therapeutically active amino acid region (IV), which covalently bonds with amino/hydroxyl/thoi groups on blood components to form a peptidase stabilised therapeutic peptide composed of 3-50 amino acids. (I) are useful for modifying therapeutic peptides e.g. hormones, growth factors and neurotransmitters, to protect them from peptidase activity in vivo for the treatment of various disorders. Endogenous therapeutic peptides are not suitable as drug candidates as they require frequent administration due to rapid degradation by peptidases in the body.
                                                                                                                                                                                                                                                                                                                                                                   Modifying and attaching therapeutic peptides to albumin prevents peptidase degradation, useful for increasing length of in vivo activity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Somatostatin analogue; syndrome X of Reaven; hyperinsulinaemia syndrome
diazoxide; cyclothiazide; metformin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Modifying and attaching therapoutic peptides to albumin prevents or reduces the action of peptidases to increase length of activity (half. life) and specificity as bonding to large molecules decreases intracellular uptake and interference with physiological processes. AAB90829 to AAB92441 represent peptides which can be used in the exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present invention describes a modified therapeutic peptide (I) comprising a therapeutically active amino acid region (III) and a reactive group (II) (e.g. succinimidyl and maletand provent.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Somatostatin analogue for the treatment of syndrome X of Reaven
                                                                                                                                                                                                                                              Bridon DP, Ezrin AM, Milner PG, Holmes DL, Thibaudeau K;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 105, DB 22; Length 25;
Pred. No. 1e-08;
2; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure; Page 252; 733pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAW51859 standard; peptide; 28 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    5 LPPRERKAGCKNFYWKGFTSC 25
                                                   99US-0134406.
99US-0153406.
99US-0159783.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Match 71.4%;
Local Similarity 81.0%;
es 17; Conservative
17-MAY-2000; 2000WO-US13576.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              97WO-IL00301.
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                                                                                                                                                                                   (CONJ-) CONJUCHEM INC.
                                                                                                                                                                                                                                                                                                         WPI; 2001-112059/12.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 25 AA;
                                                   17-MAY-1999;
10-SEP-1999;
15-OCT-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (COHE/) COHEN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO9810786-A2.
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12-SEP-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                19-MAR-1998.
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Length 28;

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The present invention describes immortalized neuroendocrine cells (A) that stably secrete a polypeptide hormone (I) contain an expression negion that includes a transgene (TG), linked to a promoter functional in enkaryctic cells, such that expression of TG increases sensitivity of the cells to a modulator of (I) secretion. (A) are specifically used, by transplantation, to treat diabetes or hypoglycaenia (especially where associated with insulin therapy) but more generally are used to express, in tvo, a wide range of therapeutic hormones, enzymes, amidated proteins and growth factors. Also engineered neuroendocrine cells are used to identify new therapeutic agents or drug targets (A) have a stable phenotype and particularly inducible julcagon secretion and glucose counter-requisatory capacities, i.e. they balance the hyperglycaemic effects of beta-cell loss and the hypoglycaemic
                                                                 but can also be used in cases of other neuroendocrine-based disorders such as Parkinson's disease, athyrotic cretinism and addison's disease. The method uses engineered, immortalised cells that are available in large amounts, with a stable and predictable phenotype. They allow screening to be performed in vivo. The present sequence represents a sometostatin peptide hormone SS-28 used in an example from the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New immortalized neuroendocrine cells that stably secrete polypeptide, particularly used to treat diabetes and hypoglycaemia
        o vivo or in vitro. Specifically the diabetes by regulation of insulin
                                                                                                                                                                                                                                                                                                                                                                                      Score 105; DB 20; Length & Pred. No. 1.1e-08; Pred. ......has 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Glucagon-like peptide I receptor; GLP-1; regulation; neuroendocrine cell line; glycaemic sensing mechanism genetic engineering; hypoglycaemia; dlabetes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 8; Page 148; 318pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Peptide hormone somatostatin SS-28.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          5 LPPRERKAGCKNFYWKGFTSC 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   8 MAPRERKAGCKNFFWKTFTSC 28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAY24239 standard; peptide; 28 AA
                                                                                                                                                                                                                                                                                                                                                                                                                  71.4%;
81.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   98US-0087848.
98US-0071193.
98US-0071209.
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98US-0087821
                                     or preventing
be used in cas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    15-SEP-1999 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity 81.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Clark SA, Thigpen AE;
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polypeptides from of for treating or prebut can also be use such as Parkinson's
                                                                                                                                                                                                                                                                                                                                                     28 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             11-JAN-1999;
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03-JUN-1998;
                                                                                                                                                                                                                                                                                   invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAY24239;
                                                                                                                                                                                                                                                                                                                                                         Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 13
AAY24239
    88888888888
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ö
                                                                              The present sequence is a mouse somatostatin (SS-28) hormone which was found to inhibit the release of growth hormone, thyroid stimulating hormone, insulin and glucagon. In addition, SS-28 and its analogue octrectide may inhibit growth of some tumours. The hormone was used to study its effect on glucose-stimulated insulin secretion in high expressing and non-expressing clones of mouses somatostatin receptor, type V gene (SSTRV). The insulin secretion was highly inhibited in the high expressing clone as compared to the non-expressing clone because the high expressing clone as compared to the non-expressing clone somatostatin. The hormone effectively inhibited insulin secretion in the absence of glucose. The somatostatin receptor can be introduced in cellines, used in cell-based delivery of insulin for treating diabetes, for precise regulation of insulin release.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present invention describes a method for identifying modulators (I) of secretory function by treating an immortalised cell, having a stable secretory function, with a test compound and detecting any change in secretion caused by the compound. (I) are used to confrol secretion of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Glucagon-like peptide I receptor; GLP-1 receptor; drug screening; secretory function; immortalised neuroendocrine secretory cell; regulation; diabetes; insulin secretion; neuroendocrine-based disorder; Parkinson's disease; athyrotic cretinism; Addison's disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                71.4%; Score 105; DB 20; Length 28; 81.0%; Pred. No. 1.1e-08; tive 2; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 8; Page 153-154; 309pp; English.
                Example 8; Page 143; 312pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Somatostatin peptide hormone SS-28.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 5 LPPRERKAGCKNFYWKGFTSC 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Ş
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAY24384 standard; peptide; 28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           98US-0071193.
98US-0071209.
98US-0072556.
98US-0087821.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  20-SEP-1999 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     similarity 81.0
17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (BETA-) BETAGENE INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Clark SA, Quaade C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 1999-430454/36.
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Mus musculus. WO9935495-A2.

AAY24384;

RESULT 12 AAY2438

Query Match Best Local S Matches 17

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Sequence

03-JUN-1998; 12-JAN-1998; 12-JAN-1998; 12-JAN-1998; 03-JUN-1998;

11-JAN-1999; 15-JUL-1999

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The present invention describes a modified therapeutic peptide (I) comprising a therapeutically active amino acid region (III) and a reactive group (II) (e.g. succinimidy) and maleimide groups) attached to a less therapeutically active amino acid region (IV), which covalently bonds with amino/hydroxyl/thiol groups on blood components to form a peptidase stabilised therapeutic peptide composed of 3-50 amino acids. (I) are useful for modifying therapeutic peptides e.g. hormones, growth factors and neurotransmitters, to protect them from peptidase activity in vivo for the treatment of various disorders. Endogenous therapeutic peptides are not suitable as drug candidates as they require frequent administration due to rapid degradation by peptidases in the body. Modifying and attaching therapeutic peptides to albumin prevents or reduces the action of peptidases to increase length of activity (half life and specificity as bonding to large molecules. decreases in the action of peptidases to large molecules. decreases in the action of peptidases with physiological processes. AMB90829 to AMB92441 represent peptides which can be used in the
effects of administered insulin. Since they are of human origin, they are less likely to suffer immune rejection than xenografts. The present sequence represents the peptide hormone somatostatin SS-28 used in an example from the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Modifying and attaching therapeutic peptides to albumin prevents peptidase degradation, useful for increasing length of in vivo activity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Protection; endogenous therapeutic peptide; peptidase; conjugation; blood component; modification; succinimidyl; maleimido group; amino hydroxyl; thiol; hormone; growth factor; neurotransmitter.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Bridon DP, Ezrin AM, Milner PG, Holmes DL, Thibaudeau K;
                                                                                                                                                                                                                                                    Watch 71.4%; Score 105; DB 20; Length 28; Local Similarity 81.0%; Pred. No. 1.1e-08; les 17; Conservative 2; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Somatostatin related peptide SEQ ID NO:192.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAB90829 to AAB92441 represent peptides exemplification of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; Page 252; 733pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                               5 LPPRERKAGCKNFYWKGFTSC 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAB91018 standard; Peptide; 28 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               17-MAY-2000; 2000WO-US13576
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      99US-0134406
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990S-0159783
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                                                                                                                                                                                 28 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             17-MAY-1999;
10-SEP-1999;
15-OCT-1999;
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Synthetic.
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                                                                                                                                                                                 Sequence
                                                                                                                                                                                                                                                             Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        rights imminodeficiency syndrome (AIDS) and various rights (e.g. epilepsy, Alzheimer's disease and asses and somatostatin antagonists are effective in the conditions. Nucleic acids encoding the polypeptides are erapy and fusion peptides can be targeted to neoplasms ses, inhibiting the release of their secretory products. Tesents the mammalian somatostatin 28 (SS-28) peptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Rainbow trout; somatostatin; preprosomatostatin; hypersecretion; PPSS-I; PPSS-II'; PPSS-II'; endocrine tumour; pituitary gland; glucagonoma; AIDS; gastroenteropancreatic tissue; acromegaly; gastrinoma; diabetes mellitus; carcinoid syndrome; cell proliferation; apoptosis; growth hormone; SS-28; glucagon; acquired immunodeficiency syndrome; neurological disorder; HIV; epilemer's disease; meturological disorder; HIV; epplease; Alchemer's disease; meturological disorder; anti-human immunodeficiency virus; osteopathic; anticonvulsant.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       secretion is
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for treating diabetes mellitus, acromegaly, gastrinoma, acquired
immunodeficiency syndrome and neurological disorders
                                                                                                                                                       ö
                                                                      Length 28;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      in the treatment of diabetes mellitus via inhibition o
and glucagon. In addition, dysfunctional somatostatin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mammalian somatostatin 28 (SS-28) peptide sequence.
                                                                  Query Match 71.4%; Score 105; DB 22; 3 Best Local Similarity 81.0%; Pred. No. 1.1e-08; Matches 17; Conservative 2; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sheridan MA, Moore CA, Kittelson JD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAU07669 standard; Peptide; 28 AA.
                                                                                                                                                                                                                         5 LPPRERKAGCKNFYWKGFTSC 25
                                                                                                                                                                                                                                                                                       8 MAPRERKAGCKNFFWKTFTSC 28
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useful in gene therapy and
and their metastases, inhib
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            04-DEC-2001 (first entry)
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28 AA;
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Sequence
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•	Score 105; DB 22; Length 28; Pred: No. 1.1e-08;	les 2; Indels 0;		
	ø ø		GFTSC 25	
28 AA;	Query Match 71.48; Best Local Similarity 81.08;	Matches 17; Conservative	5 LPPRERKAGCKNFYWKGFTSC 25	
SQ Sequence	Query Match Best Local S	tches 17	S.	
ÖS	96	Ma	à	ź

Search completed: August 13, 2003, 14:50:59 Job time : 46.7949 secs

Sequence 26, Sequence 2, Sequence 26 Sequence 38 Sequence 38

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APPLICANT SUCCLIFFE, Gregor J.
APPLICANT SUCCLIFF, Gregor J.
de Lecea, Luis
Siggins, George R.
Henriksen, Steven J.
TILE OF INVENTION: CORTISPATIN: NEUROPEPTIDES,
COMPOSITIONS AND METHODS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 3, Application US/09766396
Sequence 0. US2002001455A1
GENERAL INFORMATION:
APPLICANT: Sutcliffe, Gregor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           to have a being printed,
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134.451 Million cell updates/sec
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(egn2_6/ptodata/2/pubpaa/US07_PUBCOMB.pep:*

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                                                                                                                                                                                                                      ; Search time 24.359
GenCore version 5.1.6
Copyright (c) 1993 - 2003 Compugen Ltd.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         492763 seqs, 131003257 residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                      SVDNLPPRERKAGCKNFYWKGFTSC 25
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Maximum Match 100%
Listing first 45 summaries
                                                                                                                                                         using sw model
                                                                                                                                                                                                                                    August 13, 2003, 14:49:47
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Gapop 10.0 , Gapext 0.5
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147
1 SVDNLPPRERKAGCKNFY
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Maximum DB seq length: 200000000
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Perfect score:
Sequence:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Scoring table:
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                                                                                                                                                                                                                                                  Run on:
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NUMBER OF SEQUENCES:

ADDRESSE:
ADDRESSE:
THE SCRIPPS RESEARCH INSTITUTE
STREET: 10666 No. US20020013456A1th TOTTEY Pines Road, TPC-8
STREET: 10666 No. US20020013456A1th TOTTEY Pines Road, TPC-8
CITY: La Jolla
STREET: 10666 No. US20020013456A1th TOTTEY Pines Road, TPC-8
CONFINE EAL STREET: 10666 No. US20020013456A1th TOTTEY DESCRIPTION TOTE
COMPUTER: PLOPPY disk
COMPUTER: PLOPPY disk
COMPUTER: PROPER PC. COMPATIDE
COMPUTER: PENCHON PC. DOS/NS-DOS
SOFTWARE: PENCHON NOWBER: US/09/766,396
FILING DATE: 18-Jan-2001
FILING DATE: 18-Jan-2001
FILING DATE: CURROWNPRIOR PEDICATION NOWBER: 08/857,389
FILING DATE: CURROWNATTORNEY/AGENT INFORMATION:
NAME: Schmonsees, William
REFERENCE/DOCKET NUMBER: 31,796
REFERENCE/DOCKET NUMBER: 31,796
REFERENCE/DOCKET NUMBER: 31,796
REFERENCE/DOCKET NUMBER: 31,796
REFERENCE/DOCKET NUMBER: 32908-0002
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION 1415) 324-7041

Sequence 64, Appl Sequence 52, Appl Sequence 2, Appl Sequence 2, Appl Sequence 16, Appl Sequence 7, Appl Sequence 7, Appl Sequence 6, Appl Sequence 7, Appl Sequence 6, Appl Seq

US-10-060-375-3 US-09-280-060-53 US-09-101-487-53 US-09-316-505-2 US-10-224-640-2 US-10-224-640-2 US-10-221-703-18 US-10-101-487-37 US-10-062-375-7 Sequence

Description

Length I

Score

90 MAPRERKAGCKNFFWKTFTSC 110

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DNAS ENCODING NEW FUSION PROTEINS AND PROCESSES FOR PREPARING USEFUL POLYPEPTIDES THROUGH EXPRESSION OF THE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TOMPKINS, CHRISTOPHER K.
WAGGONER JR., DAVID W.
VENTION: RECOMBINANT PRODUCTION OF POLYANIONIC POLYMERS AND USES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; FEATURE:
; OTHER INFORMATION: Description of Artificial Sequence: Synthetic fusion
; OTHER INFORMATION: protein
US-10-101-487-53
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                OTHER INFORMATION: Description of Artificial Sequence: Designated is COTHER INFORMATION: an amino acid sequence of OTHER INFORMATION: MWPSp-MWPmp20-(His)6-EGF-TEV-Somatostatin 28 US-09-280-030-64
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 105; DB 9;
Pred. No. 6.9e-08;
2; Mismatches 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 83; DB 14;
Pred. No. 0.00017;
2; Mismatches 2
                                                                                                                                                                                                                                                                                                                                                                                                           PLICATION NUMBER: JP10-87339/1998
LING DATE: 1998-03-31
                                                                                                                    Application US/09280030A
20010021515A1
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to. US20020169125A1
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Best Local Similarity 81.0%;
Matches 17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DAVID W.
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Best Local Similarity
Matches 13; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER: US/08/857,389
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Best Local Similarity 81.0%; Pred. No. 5.4
Matches 17; Conservative 2; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CATION NUMBER: US/10/062,375
G DATE: 30-Jan-2002
IFICATION: <Unknown>
                                                                                                            MOLECULE TYPE: protein
FRAGMENT TYPE: C-terminal
SEQUENCE DESCRIPTION: SEQ ID NO: 3:
02-09-766-396-3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Henriksen, Steven J. TITLE OF INVENTION: CORTISTATIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Sutcliffe, Gregor J.
                                                                                                                                                                                                                                                                                                                                              90 MAPRERKAGCKNFFWKTFTSC 110
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   sees, William
NUMBER: 31,7
                                                                                                                                                                                                                                                                                                                 5 LPPRERKAGCKNFYWKGFTSC 25
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EDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Siggins, George R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 3, Application US/10062375
Publication No. US20020133000A1
GENERAL INFORMATION:
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FRAGMENT TYPE: C-terminal
INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADDRESS
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INFORMATION FOR

US-10-062-375-3

ATTORNEY,

PRIOR

CURRENT

RESULT 2 US-10-062-375-3

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canley, Mark S.
NTION: LOW MOLECULAR WEIGHT PEPTIDOMIMETIC GROWTH HORMONE SECRETAGOG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; OTHER INFORMATION: Description of Artificial Sequence: Extracellular OF-10-251-703-18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 77; DB 12; Length 14;
Pred. No. 9e-05;
1; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              REFERENCE: NEWC-215 CIP
ENT APPLICATION NUMBER: US/10/251,703
                                                                                                                                                                                                                                                                                                                                                                                       US 09/057,074
                                                                                                                Application US/10224640
No. US20030139348A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     12 AGCKNFYWKGFTSC 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 AGCKNFFWKTFTSC 14
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Best Local Similarity 85.7
Matches 12; Conservative
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Best Local Similarity 85.7
Matches 12; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ORGANISM: Homosapiens
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SOFTWARE: PatentIn
SEQ ID NO 18
LENGTH: 14
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ION: LOW MOLECULAR WEIGHT PEPTIDOMIMETIC GROWTH HORMONE SECRETAGOGUES
PORSIODS
                                                                                                                                                                                           LEUNG, DAVID W.
BERGARAN, PHILIP A.
LOFOUIST, ALAN
PIETZ, GREGORY E.
TOMPRINS, CHRISTOPHER K.
WAGGONEN JR., DAVID W.
FENTION: RECORDINANT PRODUCTION OF POLYANIONIC POLYMERS AND USES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; FEATURE:
; OTHER INFORMATION: Description of Artificial Sequence: Synthetic peptide
0S-10-101-487-52
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Ouery Match ,53.1%; Score 78; DB 14; Length 15; Best Local Similarity 80.0%; Pred. No. 6.8e-05; Matches 12; Conservative 2; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Ouery Match 52.4%; Score 77; DB 10; Length 14; Best Local Similarity 85.7%; Pred. No. 9e-05; Matches 12; Conservative 1; Mismatches 1; Indels
                                                                                                                                       Application US/10101487
5. US20020169125A1
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                      9 ERKAGCKNFYWKGFTSC 25
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: 1998-04-08
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PRIOR FILING DATE: 2001-
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Length 29; Indels

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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/062,375
FILING DATE: 30-Jan-2002
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TON NUMBER: 08/857,389
                                                                                                                                                                                                                                                                                                                                                                              Score 77;
                                                                                                                                                                                                                                                                            MOLECULE TYPE: protein
FRAGMENT TYPE: C-terminal
SEQUENCE DESCRIPTION: SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      9 PPHRDKKPCKNFFWKTFSSC 28
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tion No. US20020133000A1
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Best Local Similarity 60.0%;
Matches. 12; Conservative
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                                                                                                                   TELECOMMUNICATION
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                                                                                                                                                                                                                                                                        PIETZ, GREGORI E.
TOMENIN, CHRISTOOHER K.
MAGGONER JR., DAVID W.
ENTION: RECOMBINANT PRODUCTION OF POLYANIONIC POLYMERS AND USES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; OTHER INFORMATION: Description of Artificial Sequence: Recognition US-10-101-487-37
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STREET: 10666 No. US20020013456Alth Torrey Pines Road
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Stitcliffe, Gregor J.
APPLICANT: Stitcliffe, Gregor J.
Signing, George R.
Henriksen, Steven J.
                                                                                                                                                      Application US/10101487
No. US20020169125A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 7, Application US/09766396
Patent No. US20020013456A1
                                                                                                                                                                                         BERCHAN, PHILIP A.
OPQUIST, ALAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ORGANISM: Artificial Sequence
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Best Local Similarity 85.7%;
Matches 12; Conservative
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                  12 AGCKNFYWKGFTSC 25
                                     1 AGCKNFFWKTFTSC 14
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PRIOR FILING DATE: 2001-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQ ID
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RESULT 12 nc-09-766-396-6

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ADDRESSEE: THE SCRIPPS RESEARCH INSTITUTE
STREET: 10666 No. US20020133000Alth Torrey Pines Road, TPC-8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        "WT APPLICATION DATA:
APPLICATION NUMBER: US/10/062,375
FILING DATE: 30-Jan-2002
CLASSIFICATION: <a href="https://doi.org/10/10/2012/">CLASSIFICATION: <a href="https://doi.org/10/2012/">CLASSIFICATION: <a href="https://doi.org/10/2012/">https://doi.org/10/2012/">CLASSIFICATION: <a href="https://doi.org/10/2012/">https://doi.org/10/2012/">https://doi.org/10/2012
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US/08/857,389
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MOLECULE TYPE: protein
\FRAGERIT TYPE: C-terminal
SEQUENCE DESCRIPTION: SEQ ID NO: 6:
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No. US20020013456Al
RAL INFORMATION:
APPLICANT: Sutcliffe, Gregor J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    6 PPRERKAGCKNFYWKGFTSC 25
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Siggins, George F
Henriksen, Steven
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                                                                                                                                                                                                                                       TITLE OF INVENTION: CORT
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                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES:
CORRESPONDENCE ADDRES
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                                 GENERAL INFORMATION:
APPLICANT: Sutc
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                                                                                                                      Ouery Match 52.4%; Score 77; DB 14; Length 29; Best Local Similarity 60.0%; Pred. No. 0.00019; Matches 12; Conservative 2; Mismatches 6; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
COTMARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US/09/766,396
FILING DATE: 18-Jan-2001
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2; Mismatches
SEQUENCE DESCRIPTION: SEQ ID NO: 7: US-10-062-375-7
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FRAGMENT TYPE: C-terminal

SEQUENCE DESCRIPTION: SEQ ID NO: 6:
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Sutcliffe, Gregor J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     de Lecea, Luis
Siggins, George R.
Henriksen, Steven J
TITLE OF INVENTION: CORTISTATI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                           6 PPRERKAGCKNFYWKGFTSC 25
                                                                                                                                                                                                                                                                                                                                               9 PPHRDKKPCKNFFWKTFSSC 28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             pplication US/09766396
20020013456A1
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Publication No. US20020133000A1
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SEQUENCE CE
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Query Match Best Loc Matches

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SYSTEM: PC-DOS/MS-DOS
Patentin Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                Score 77; DB 9;
Pred. No. 0.00071;
2; Mismatches 6
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60.0%; Pred. No. 0.00071;
tive 2; Mismatches 6
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E: 1488.043003
CATION UNMER: US/10/335,125
S DATE: 2003-01-02
                 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/766,396
                                                                                                                                                                                                                                                                                MOLECULE TYPE: protein
SEQUENCE DESCRIPTION: SEQ ID NO: 2:
"US-09-766-396-2
                                                                                        APPLICATION NUMBER: 08/857,389
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US/09/775,827A
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R: US_60/037,386
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12-31
                                                                    <Unknown>
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                                                                                                                                                                                                                                                                                                                                                                                                  Application US/10335125
No. US20030148355A1
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                                                                                                                                                                                                                                                                                                                                              Query Match 52.4%;
Best Local Similarity 60.0%;
Matches 12; Conservative
                                                                                 PRIOR APPLICATION DATA:
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Best Local Similarity 60.0
Matches 12; Conservative
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; ORGANISM: Rat Cortistatin
US-10-335-125-3
                                                                                                                                                                                                                     INFORMATION FOR SEQ I
SEQUENCE CHARACT
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LENGTH: 112
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Search completed: August 13, 2003, 14:53:43

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Periect score: 147
Sequence: 1 SYDNLPPRERKAGCKNFYWKGFTSC 25
___Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 283308 seqs, 96168682 residues

Total number of hits satisfying chosen parameters:

283308

Minimum DB seq length: 0
Maximum DB seq length: 200000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
Database .: PIR_76:*
1: pir1:*
3: pir2:*
3: pir3:*
4: pir4:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

		Description	somatostatin-25 -	somatostatin II pr	somatostatin II pr	somatostatin II pr	somatostatin II pr	somatostatin-25 -	somatostatin-14 pr	preprosomatostatin	somatostatin-14 pr	somatostatin precu	somatostatin I pre	somatostatin-28 -	somatostatin I pre	somatostatin I pre	somatostatin I pre	somatostatin precu	somatostatin precu	somatostatin precu	somatostatin-relat	somatostatin, panc	somatostatin-14 [P	somatostatin - sli	somatostatin I - c	somatostatin I - E	somatostatin I - s	cortistatin precur	somatostatin - spo	~	hypothetical prote
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			i -				•																								
		ID	A60842	RIAFS2	800169	IS1064	800166	B60840	RIIDS1	I50798	JC6166	S20630	RIAFSI	A61322	RIPGS	RIHUS1	A28968	RIBOS1	RIRTS1	RIMSS1	A32271	A32000	JC6167	C60414	B60842	A60840	S00172	S67489	A60622	JC5414	F86157
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		Match Length DB	25	125	73	115	74	25	114	114	115	116	121	28	92	116	116	116	116	116	34	37	103	14	14	14	14	112	14	105	316
م ه	Query	Match	100.0	93.9	91.2	91.2	84.0	77.6	72.8	72.8	72.8	72.8	72.8	71.4	71.4	71.4	71.4	71.4	71.4	71.4	69.4	63.9	58.2	52.4	52.4	52.4	52.4	52.4	49.0	44.2	35.4
		Score	147	138	134	134	123.5	114	101	107	107	107	101	105	105	105	105	105	105	105	102	94	85.5	77	77	77	77	77	72	65	22
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-		סטכנודה ז
		A60842
		coho salmon
		N; Alternate names: somatostatin II precursor
	•	C;Species: Oncorrypromis Kishuch (coho salmon) C:Date: 10-sen-1903 #sequience revision 30-sen-1903 #fext change 07-May-1999
7		
		isetskaya, E.M.; Pollock, H.G.;
	•	A;TLLE: Characterization of Lono salmon (Oncorny)renus Kisuren) islet somatostatins A;Reference number: A60842; WUID:87055212; PMTD:2877919
		A; Accession: A60842
		A; Molecule type: protein
		A; Nestudes: 1-20 - KFLL> A: Access for: C60842
		A; Molecule type: protein
		(PL2>
		A; Note: this form, sometostatin II, was not sequenced directly but rather deduced I)
		C.Subellamily: Solutioscalin C.Kavwords: hormone: nancreatic felet
		c.n.g.m.um.m.m.r.p.m.r.m.r.r.r.r.r.r.r.r.r.r.r.r.
		s experimental
		Query Match 100.0%; Score 14/; DB 2; Length 25;
		vative O Mismatch
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		T SADARE EXERCISES THE TANGE TO
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		RESULT 2
		Sometostatin II precursor - American quosefish
		lus americanus (American goosefish)
		C;Date: 13-Mar-1981 #sequence_revision 31-Mar-1981 #text_change 28-May-1999
		C./ACCESSION: 59220; ARYCOO; ALISTO, A
-		;
		A; Title: Cloning and sequence analysis of cDNAs encoding two distinct somatostatin [
		A:Reference number: A93236; MUID:81052423; PMID:6107860
		A;Accesson: b93429 A;Molecule type: mRNA
		A; Residues: 1-125 <hob></hob>
		V00641
		A:Experimental Source: 1Siet tissue (endocrine pancreas) R:soless II Nos R D
		Proc. Natl. Acad. Sci. U.S.A. 82, 277-281, 1985
		an anglerfi
•		A; ACCESSION: AV&USO

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Length 115;

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A; Molecule type: protein
A; Molecule type: protein
A; Molecule type: protein
A; Molecule three peptides which probably originate from a common precursor, were isolate
A; Note: the source is designated as Cottus scorpius
R; Cutfield, S.M.; Carne, A.; Cutfield, J.F.
FEBS Lett. 214, 57-61, 1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Tr. J. Blochem. 168, 647-652, 1987

Title: Structural characterization of peptides derived from prosomatostatins I and Reference number: S00166; MUID:88029486; PMID:2889597
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A;Title: The amino-acid sequences of sculpin islet somatostatin-28 and peptide YY.
A;Reference number: A91376; MUID:87190954; PMID:2883025
A;Recession: A26993
A;Molecule type: protein
A;Residues: 47-74 <CUTY>
C;Superfamily: somatostatin
C;Reywords: Glycoprotein; neuropeptide; pancreatic islet
F;1-12/Product: peptide S1 #status experimental <PS1>
F;13-4/Product: peptide S2 #status experimental <PS2>
F;13-74/Product: peptide S2 #status experimental <PS3>
F;13-74/Product: peptide S3 #status experimental <PS3>
F;13-74/Product: peptide S4 #status experimental 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 somatostatin II precursor - shorthorn sculpin (tentative sequence) (fragments) C.Species: Myoxocephalus scorpius (shorthorn sculpin, daddy sculpin) C.Date: 07-Sep-1990 #sequence_revision 26-Jan-1996 #text_change 31-Mar-2000 C.Accession: S00166; A26993
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Similarity 95.7%; Pred. No. 1
22; Conservative 1; Mismatch
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1 Similarity 82.18;
23; Conservative
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Best Local Similarity 84.0.
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Molecule type: protein
Residues: 99-125 <- SPET>
Residues: 98-125 <- Residues: 98-125 
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*Residues: 1-76, DV. 79-89, 'G', 91-125 <AND>
*Residues: 1-76, DV. 79-89, 'G', 91-125 <AND>
*Superfamily: somatostatin
iKeywords: hydroxylysine; neuropeptide; pyroglutamic acid
il-24 Domain: signal sequence #status experimental <SIG>
ig-25-97/Domain: propeptide #status experimental <PRO>
ig-125/Product: somatostatin II #status experimental <AND>
ig-125/Modified site: pyrrolidone carboxylic acid (Gln) (in mature form) #status experimental
il14-125/Disulide bonds: #status experimental
il120/Modified site: hydroxylysine (Lys) #status experimental
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conlon, J.M.; Davis, M.S.; Falkmer, S.; Thim, L.
T. Blochem. 186, 647-652, 1987.
Title: Structural characterization of peptides derived from prosomatostatins I and II
Reference number: S00166; MUID:88029486; PMID:2889597
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Gen. Comp. Endocrinol. 98, 253-261, 1995
A;Title: Isolation and characterization of a cDNA encoding for preprosomatostatin contain
A;Reference number: I51064; MUID:95354921; PMID:7628684
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  - European flounder (tentative sequence) (fragments)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            somatostatin II precursor - Buropean flounder (tentative sequence) (fragment
C:Species: Platichthys flesus (European flounder)
C:Date: 07-Sep-1990 #sequence_revision 07-Sep-1990 #text_change 31-Mar-2000
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C:Species: Oncorhynchus mykiss (rainbow trout)
C:Date: 13-Sep-1996 #sequence_revision 13-Sep-1996 #text_change 16-Jul-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ö
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95.7%; Pred. No. 7.1e-13;
tive 1; Mismatches 0
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A;Status: preliminary; translated from GB/EMBL/DDBJ
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Pred. No. 3e-13;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity 92.0%; Pred. No. 3e-1
Matches 23; Conservative 1; Mismatches
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Best Local Similarity 95.77
Watches 22; Conservative
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Moore, C.A.; Kitt
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Cincession: B66840

Riconion, J.M.; Deacon, C.F.; Hazon, N.; Henderson, I.W.; Thim, L.

Riconion, J.M.; Deacon, C.F.; Hazon, N.; Henderson, I.W.; Thim, L.

Gen, Comp. Endocrinol. 72, 181-189, 1989, 1989, 1981

A; Feference number: A60840; MUID: 89065329; PMID: 2904391

A; Reference number: A60840; MUID: 89065329; PMID: 2904391

A; Reference number: A60840; MUID: 89065329; PMID: 2904391

A; Reference number: A60840; MUID: 89065329; PMID: 2904391

A; Residual type: protein

A; Residual type: protein type: pr
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C;Species: Anguilla anguilla (European eel)
C;Date: 30-Sep-1993 #sequence_revision 30-Sep-1993 #text_change 20-Mar-1998
C;Accession: B60840
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Score 123.5; DB 2;
Pred. No. 2.6e-11;
1; Mismatches 1;
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Pred. No. 2.4e-10;
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somatostatin I precursor - American goosefish
%;Contains: somatostatin is Captains and Marchaelian
%;Species: Lophius americanus (American goosefish)
%;Species: Lophius americanus (American goosefish)
%;Species: Lophius americanus
%;Marchaelian: A32326; A33860; A91087; A01433
R;Hobart, P.; Crawford, R.; Shoi, L.; Pitter, R.; Rutter, W.J.
%;Mature 288, 137-141, 1980
A;Hitle: Clouing and sequence analysis of CDNAs encoding two distinct somatostatin pa
A;Reference number: A93236; MUID:81052423; PMID:6107860
                                                                                                                  rocettvint, H.; Lihrmann, I.; Bucharles, C.; Vieau, D.; Coulouarn, Y.; Fournier, A roc. Ratl. Acad. Sci. U.S.A. 93, 12665-12610, 1996
Title: Occurrence of two somatostatin variants in the frog brain: Characterization Reference number: JC6166; MUID:97057290; PMID:8901629
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R;Nata, K.; Kobayashi, T.; Karahashi, K.; Kato, S.; Yamamoto, H.; Yonekura, H.; Okam Submitted to the EMBL Data Library, June 1991
S;Description: Nucleotide sequence determination of chicken somatostatin precursor c A;Reference number: S20630
                                                                                                                                                                                                                                                                                                                                               Molecule type: mRNA
Residues: 1-115 CTOSS: TOSS: 01D:91890650; PIDN:AAC60093.1; PID:91890651
COOSS-references: GB:068136; NID:91890650; PIDN:AAC60093.1; PID:91890651
COMMENT: This protein acts both as a neurotransmitter/neuromodulator and a hormone
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                                                                C;Species: Rana ridibunda (laughing frog)
C;Date: 11-Apr-1997 #sequence_revision 09-May-1997 #text_change 16-Jul-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        somatostatin precursor - chicken
C;Species: Gallus gallus (chicken)
C;Date: 10-Sep-1999 #sequence_revision 10-Sep-1999 #text_change 10-Sep-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A:Cross-references: EMBL:X60191; NID:962985; PIDN:CAA42747.1; PID:962986 C;Superfamily: somatostatin
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85.7%; Pred. No. 1.1e-08;
1ve 1; Mismatches 2; Indels
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Best Local Similarity 85.7
Matches 18; Conservative
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Best Local Similarity 85.7
Matches 18; Conservative
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C;Superfamily: somatostatin
C;Keywords: brain; hormone
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A: Residues: 1-116 <NAT>
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                                                                          C; Species: Ictalurus punctatus (channel catfish)
C;Date: 30-0301-1980 Seequence_revision 31-Dec-1993 #text_change 18-Jun-1999
C;Date: 30-0302; A93897; A92334; A01435
R;Minth, C.D.; Taylor, W.L.; Magazin, M.; Tavianini, M.A.; Collier, K.; Weith, H.L.; Dix J. Biol. Cham. 257, 10372-10377, 1982
A;Itle: The structure of cloned DNA complementary to catfish pancreatic somatostatin-14
A;Reference number: S00292; MUID:82265698; PMID:6179939
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Molecule type: mRNA

*Residues: 82-108 (-700)

*Andrews, P.C.; Dixon, J.E.

*Andrews, P.C.; Baron, J.E.

*Andrews, P.C.; Baron, J.E.

*Andrews, P.C.; Baron, J.E.

*Andrews, P.C.; Server, J.E.

*Andrews, P.C.; Server, J.E.

*Andrews, P.C.; Paron, J.E.

*Andrews, P.C.; Paron, J.E.

*Andrews, P.C.; Paron, J.E.

**Andrews, P.C.; Paron, J.E.
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R.Taylor, W. L.; Collier, K.J.; Deschenes, R.J.; Wetth, H.L.; Dixon, J.E.
Proc. Natl. Acad. Sci. U.S.A. 78, 6684-6689, 1981
A.Title: Sequence analysis of a cDNA coding for a pancreatic precursor to somatostatin.
A.Reference number: A93897; MUID:82082515; PMID:6171821
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Species: Ictalurus punctatus (channel catfish)
Date: 13-Sep-1996 #sequence_revision 13-Sep-1996 #text_change 16-Jul-1999
Accession: 150798
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A; Residues: 1-114 cDIX>
A; Residues: 1-114 cDIX>
A; Cross-teferences: GB:M25903; NID:g213339; PIDN:AAA49339.1; PID:g213340
C; Superfamily: somatostatin
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Pred. No. 1.1e-08;
2; Indels
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752-100/Domain: propeptide #status predicted <PRO>
F:101-114/Product: somatostatin-14 #status experimental <MAT>
F:103-114/Disulfide bonds: #status experimental <MAT>
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Adv. Exp. Med. Biol. 188, 19-29, 1985
Affitle: Sometostatins of the channel caffish.
A; Reference number: 150798; MUID:85303576; PMID:2863931
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 107; DB 1;
Pred. No. 1.1e-08;
1; Mismatches 2;
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- channel catfish
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Local Similarity 85.7%;
Les 18; Conservative 1
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Best Local Similarity 85.7
Matches 18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             sidues: 101-114 <AND>
perfamily: somatostatin
sywords: neuropeptide
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Best Local S
Matches 18
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Jalternate names: prosomatostatin (SC-28) (SC-
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A; Mesidues: 65-92 - CRAA.
A; Residues: 65-92 - CRAA.
A; Experimental source: intestine
R; Schally, A.V.; Huang, W.Y.; Chang, R.C.C.; Arimura, A.; Redding, T.W.; Millar, R.P.
Proc. Natl. Acad. Sci. U.S.A. 77, 4489-4493, 1980
A; Title: Isolation and structure of pro-somatostatin: a putative somatostatin precur:
A; Reference number: A93854; MUID:81054799; PMID:6107906
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Schmidt, W.E.; Mutt, V.; Kratzin, H.; Carlquist, M.; Conlon, J.M.; Creutzfeldt, W. 1921, 141-146, 1985
Statt. 1921, 141-146, 1985
Fible: Isolation and characterization of prossl-32, a peptide derived from the N-tribeference number: A24222; MUID:86030691; PMID:2865169
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     adayrol, L.; Jornvall, H.; Mutt, V.; Ribet, A.
Lett. 109, 55-58, 1980
tle: N-terminally extended somatostatin: the primary structure of somatostatin-2
ference number: A91273; WUID:80113258; PMID:7353633
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sidues: 65-28 - SCH>
perimental source: hypothalamus
parimental source: hypothalamus
shally, A.V.; Dupont, A.; Arimura, A.; Redding, T.W.; Nishi, N.; Linthicum, G.L.;
shallstry 15, 509-314, 1976
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saidues: 79-92 «SC2»
xperimental source: hypothalamus
ersani, M.; Johnsen, A.H.; Holst, J.J.
S. Ett. 279, 237-239. 1991
11Le: Oxidation,reduction explains heterogeneity of pancreatic somatostatin.
eference number: $13616; MUID:91160722; PMID:1672110
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Title: Isolation and structure of somatostatin from porcine hypothalami
Reference number: A90398; MUID:76186331; PMID:1252409
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81.0%; Pred. No. 1.8e-08;
ative 2; Mismatches 2;
           (fragment)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity 81.0
Matches 17; Conservative
                 somatostatin I precursor - pig
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Superfamily: somatostatin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A; Molecule type: protein
A; Residues: 1-64 <BER>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   dues: 1-32 <SC3>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Molecule type: protein
Residues: 79-92 <BE2>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Accession: A24222
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Accession: S13616
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Molecule type:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Nolecule type: protein.
Residues: 1-28 cSPI.
Burgus, R.; Ling, N.; Butcher, M.; Guillemin, R.
coc. Natl. Acad. Scil. U.S.A. 70, 684-688, 1973.
Title: Primary structure of somatostatin a hypothalamic peptide that inhibits the secrence number: A61344; MUID: 72209562; PMID:4514982
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ;Noe, B.D.; Spiess, J.; Rivier, J.B.; Vale, W. adocrinology 105, 1410-1415, 1979.
;Title: Isolation and characterization of somatostatin from anglerfish pancreatic islet; Reference number: A91087; MUID:80046482; PMID:387385
Cross-references: GB:V00640; GB:J00946; NID:g64028; PIDN:CAA23986.1; PID:g64029
G60cdman, R.H.; Jacobs, J.W.; Chin, W.W.; Lund, P.K.; Dee, P.C.; Habener, J.F.
roc. Natl. Acad. Sci. U.S.A. 77, 5869-5873, 1980
Title: Nucleotide sequence of a cloned structural gene coding for a precursor of panci
Reference number: A93860; MUID:81077276; PMID:6108560
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         :Title: Isolation and sequence analysis of a somatostatin-like polypeptide from ovine Reference number: A61322; MUID:81184502; PMID:7225368
                                                                                                                                                                                                                                                                                                                                                                                                Experimental source: islet tissue (endocrine pancreas).
Goodman, R.H.; Jacobs, J.W.; Chin, W.W.; Lund, P.K.; Dee, P.C.; Habener, J.F.
roc. Natl. Acad. Sci. U. S.A. 79, 1682, 1982
Reference number: A93905
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Species: Ovis orientalis aries, Ovis ammon aries (domestic sheep)
Date: 17-Jul-1994 #sequence_revision 17-Jul-1994 #text_change 07-May-1999
Accession: A61322; A61344
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               # Match 72.8%; Score 107; DB 1; Length 121; Local Similarity 85.7%; Pred No. 1.2e-08; Andrews 18; Conservative 1; Mismatches 2; Indels hes 18; Conservative 1; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (108-121/Product: somatostatin I *status experimental <MAT>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        C:Reywords: neuropeptide
F:)-128/Product: somatostatin-28 #status experimental <528>
F:)5-28/Product: somatostatin-14 #status experimental <514>
F:)7-28/Pisulfide bonds: #status experimental
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               4/Domain: signal sequence #status predicted <SIG>
                                                                                                                                                                                                                                                                                                                                                       ,22-82,'E',84-121 <GOO>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          101 LAPRERKAGCKNFFWKTFTSC 121
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     5 LPPRERKAGCKNFYWKGFTSC 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  5 LPPRERKAGCKNFYWKGFTSC 25
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Spiess, J.; Villarreal, J.; Vale, W. cchemistry 20, 1982-1988, 1981
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Contents: annotation; erratum
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity 81.09
Matches 17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Superfamily: somatostatin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Superfamily: somatostatin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Contains: somatostatin-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Molecule type: protein
Residues: 108-121 <NOE>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Molecule type: protein
Residues: 15-28 <BUR>
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                                                                                                                                                                                                                                                          Accession: A93860
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0; Gaps

N; Alternate names: preprosomatostatin somatostatin I precursor - human

RIPGS

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PID:9342299

GB.M19318; NID:9342298; PIDN:AAA36908.1; atin inhibits the release of somatotropin.

Length 116; Indels

; DB 1; 2:3e-08;

Score 105; DB Pred: No. 2:3e-2; Mismatches

71.4%; 81.0%;

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F.1-24/Domain: signal sequence #status predicted <SIG> F:25-89/Domain: propeptide #status predicted <PRO> F:95-89/Domain: propeptide #status predicted <PRO> F:98-116/Product: somatostatin-28 #status predicted <M28> F:103-116/Product: somatostatin-14 #status predicted <M14> F:105-116/Disulfide bonds: #status predicted
         number: A28968; MUID:88144503; PMID:2894033
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Search completed: August 13, 2003, 14:51:52 Job time : 17.0256 secs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      96 MAPRERKAGCKNFFWKTFTSC 116
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    5 LPPRERKAGCKNFYWKGFTSC 25
                                                                                                                                                                                                                                                                                                                                                                                                      Best Local Similarity 81.0
Matches 17; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      e-specific mutagenesis identifies amino acid residues critical in prohormone number: $09381; MUID:90059875; PMID:2573512
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Description: inhibits the secretion of a number of peptide hormones, including somator
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1. Acad. Sci. U.S.A. 85, 1696-1700, 1988
Phenol emulsion-enhanced DNA-driven subtractive CDNA cloning: isolation of low
                                                 #sequence_revision 17-Dec-1982 #text_change 18-Jun-1999 A01430; S09381; S50024
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ascicularis (crab-eating macaque)
#sequence_revision 31-Jan-1997 #text_change 18-Jun-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             oss-references: GB:J00306; NID:g338287; PIDN:AAA60566.1; PID:g338288
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Pancreatic somatostatinoma
G; Zollinger, L.; Nault, C.; Rholam, M.; Cohen, P.
1989
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sidues: 53-62;67-82 <ODU>
ment: Somatostatin inhibits the release of somatotropin.
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(SS-14); somatostatin 28 (SS-28)
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                                                                                                                                                                                                                                                                                                                                     Acad. Sci. U.S.A. 79, 4575-4579, 1982
man somatostatin I: sequence of the cDNA.
number: A01430; MUID:83014931; PMID:6126875
                                                                                                                                                 juence of the human somatostatin I gene.
number: A43614; MUID:84146798; PMID:6142531
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Pred. No. 2.3e-08;
2; Mismatches 2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Cross-references: GDB:119604; OMIM:182450
Map position: 3q28-3q28
Introns: 46/3
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Best Local Similarity 81.0
Matches 17; Conservative
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number: S50024;
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11-11-1-1	Gencore Version 5.1.6	. 35	47.5	32.3	.372
o) language of the comparison	(c) 1993 - 2003 Compugen Ltd.	36	47	32.0	584
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ON prototo a patenta		38	45.5	31.0	147
or process . process search, using sw model	using sw model	. 39	45	30.6	216
Pur on The Contract 12		. 40	45	30.6	994
	August 13, 2003, 14:47:17; Search time 9.61539 Seconds	41	45	30.6	1109
	(without alignments)	42	44.5	30.3	518
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Sequence: 1 SVDNLPPRERK	KAGCKNFYWKGFTSC 25				
Scoring table: BLOSUM62	u C +		· . . · ·	•	

Total number of hits satisfying chosen parameters: 127863 seqs, 47026705 residues Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Scoring table: BLOSUM62 Gapext 0.5

Searched:

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

SwissProt_41:* Database :

Pred. No. 1s the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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	,	Description	P01170 lophius ame	platicht		_	_	_			•		P33094 gallus gall	P01169 lophius ame		P26917 bos taurus	P49670 canis famil	homo s	m snm	046688 ovis aries	P19209 myxine glut		_	O9ygh3 carassius a	P21779 petromyzon		P87385 rana ridibu	0	P31885 alligator m	Q62949 rattus norv	P56469 mus musculu	0	m	_	P01172 ictalurus p
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SUMMARIES		Ð	SMS2_LOPAM	SMS2_PLAFE	SMS2_ONCMY	SMS2_MYOSC	SMS2_ORENI	SMS2_CARAU	SMS1_AMICA	SMS1_ICTPU	SMSA_CARAU	SMS1_RANRI	SMS_CHICK	SMS1_LOPAM	SMS_PIG	SMS_BOVIN	SMS_CANFA	SMS_HUMAN	SMS_MOUSE	SMS_SHEEP.	SMS_MYXGL	SMS1_PROAN	SMS2_PROAN	SMSB_CARAD	SMS_PETMA	SMS_LAMFL	- 1	SMS1_MYOSC	SMS_ALLMI	CORT_RAT	CORT_MOUSE	CORT_HUMAN		TREA EMENI	SMSZ_ICTPU
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	Query	Match	93.9	91.2	91.2	84.0	82.3	74.8	72.8	72.8	72.8	72.8	72.8	72.8	71.4	7.1.4	71.4	71.4	71.4	71.4	69.4	67.3	9.0	0.00	200	500	58.2	52.4	52.4	52.4	50.3	2.44	4.00	n (7.70
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AN P26992 homo sapien 008406 rattus norv 098410 chlamydia p P9810 chlamydia p P98810 chlamydia p P98810 chlamydia p P98868 lymmatria d G92105 r sarcoplas P3489 caprine art p36734 human papil P57421 buchnera ap P10768 homo sapien P14347 epstein-bar AENTS	update) ion update) ion update) is: [Tyr7,Gly10]somatostatin-14]. fish); (Anglerfish). fish); (Anglerfish). fish); (Anglerfish). forbilitormes; Lophildae; Lophius. Fictet R., Rutter W.J.; cDNAs encoding two distinct he endocrine pancreas of	AL SEQUENCE, AND HYDROXYLATION. NE-87308304; PubMed-288757; "translational processing of preprosomatostatin-II examined fast atom bembardment mass spectrometry."; "Chem. 262:1269212699(1987). "Chem. 262:1269212699(1987). UNCTION: SORATOSTATIN INTERFERENCE OF SOMATOTROPIN. UNCTION: SORATOSTATIN ILLAY HAVE A DIFFERENT DEGREE OF CITYITY OR A DIFFERENT TYPE OF TARGET CELL FROM SOMATOSTATIN I. IMILARITY: BELONGS: TO THE SOMATOSTATIN I. MILARITY: BELONGS: TO THE SOMATOSTATIN I. INTILARITY: BELONGS: TO THE SOMATOSTATIN I. INTILARITY: BELONGS: TO THE SOMATOSTATIN I. SOMATOSTATIN I. INSTItution as a long as its content is in no way led and this statement is not removed. Usage by and for commercial is not removed. Usage by and for commercial is not removed. Usage by and for commercial is requires a license agreement (See http://www.isb-sib.ch/announce/nd an email to licensedisb-sib.ch). PPOS IRROA250; Somatostatin. PPOS IRROA250; SOMATOSTATIN-14. EN
CMTR_HUMAN CMTR_RAT TASA_CHLIN TASA_HELPN TASA_HELPN TASA_HELPN TASA_HELPN TASA_HELPN TASA_HUMAN TASA_HUMAN TASA_HUMAN TASA_HUMAN TASA_HUMAN	PRT;] sed) sequence up sequence up sequence up contains: [n goosefis) a; Cranlate; i Teleostis crygil; Lop 07860; sis of CDN nd in the e	Y572; NG OF Prep, mass spect mass spect gy(1987). NHIBITS TIN MY. TYPE OF TA TIN IN MY. TYPE OF TA HE SOMATOS. TY
CNTR CNTR Y328 Y344 YGDK YGDK VLL POL VLL FLGD ESTD UL49	DARD; PRT; 12 1, Created) 2, Last sequence upd 2, Last sequence upd 2, Last anotation ucursor (Contains: [Tamerican goosefish) Charloan goosefish) Charloan goosefish; pterygil; Teleostei; canthopterygil; Loph ubMed-6107860; Teleostei; canthopterygil; Loph ubMed-6107860; Ce analysis of CDNAs sors found in the en	HYDROXYLATION Wed=2887572; 1 R. Dixon J. 1 R. Copyright. 2 R. Copyright. 3 R. Copyright. 3 R. Copyright. 3 R. Copyright. 3 R. Copyright. 4 R. POPTENT. 5 R. HYDRON G. 6 R. Copyright. 5 R. Copyright. 6 R. Copyright. 7 R. Cop
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372 372 584 584 370 1147 216 994 1109 518 236 236 591	STANDARD; PRT; 12 (Rel. 01, Created) (Rel. 12, Last sequence upd (Rel. 10, Last annotation un II precursor (Contains: [Tricanus (American goosefish) Metazoa; Chordata; Craniata; gli, Neopterygli; Teleostei; gli, Neopterygli; Teleostei; gli, Neopterygli; Loph Mn.A. 52423; PubMed-6107860; Crawford, R., Shen L., Pict d sequence analysis of CONAs precursors found in the en 137-141(1980).	INL SEQUENCE, AND HYDROXYLATION INE-87308304; PubMed-2887572; www F.C., Mitchols R., Dixon J. E. translational processing of py gfast atom bombardment mass sp lol. Chem. 262.12692-12699(1987 GENCTION: SOMATOSTARIN INHIBITS SUBSCELLALAR LOCATION: Secreted. MACSELLALAR COCATION: SECRETEGE STRICKLIANE SECONGS. TO THE SOMA STRICKLIANE SOMA THE
220.00.00.00.00.00.00.00.00.00.00.00.00.	STANDARD STANDARD POLITO, 091066; TOT-1010, 091066; TOT-1080 (Rel. 12, L. 16-0CT-2001 (Rel. 12, L. 16-0CT-2001 (Rel. 40,	ON TO WALL THE STATE OF THE STA
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44 4 4 CC 8 4 44.444	LD 1A LOCAM LOCAM 21-CD-1986 01-CCT-1989 01-CCT-1989 16-CCT-2001 Sometostatin Lophius amer Eukaryota	PARTILE SEDU MEDLINE-8730 MEDLINE-8730 MEDLINE-8730 MADGREWS P. C. *POST-LTARNILL USING FOR THE SENDELLU TO SIMILARY TO SIMILA
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NCBI_TaxID-8097;
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Best Local S
Matches 22
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                                                                                                                                                                                                                           01-MMY-1991 (Rel. 18, Created)
01-MMY-1991 (Rel. 18, Last sequence update)
16-0CT-1991 (Rel. 40, Last sequence update)
16-0CT-2001 (Rel. 40, Last annotation update)
Somatostatin II precursor (Contains: [Tyr21,61y24]somatostatin-28;
[Tyr7,61y10]somatostatin-14] (Fragments)
Platichthys flesus (European flounder)
Eukaryota: Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Actinoperygli; Relegostel; Euteleostel;
Acanthomorpha; Acanthopetrygli; Percomorpha;
Pleuronectoldel; Pleuronectidae; Platichthys
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            16-0CT-2001 (Rel. 40, Created)
16-0CT-2001 (Rel. 40, Last sequence update)
16-0CT-2001 (Rel. 40, Last annotation update)
Somatostatin In precursor (Contains: [Tyr21;Gly24]somatostatin-28; [Tyr7,Gly10]somatostatin-14].
                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE.
TISSUE-Pancreas;
MEDLINE-88029486; PubMed-2889597;
MEDLINE-88029486; PubMed-2889597;
Conlon J.M., Davis M.S., Falkmer S., Thim L.;
Structural characterization of peptides derived from prosomatostatins 1 and 11 isolated from the panoreatic islets of prosomatostatins 1 and 11 isolated from the panoreatic islets of percentages of teleostean fish: the deddy sculpin and the flounder.";
Suc. J. Biochem. 168:647-652(1987).
I. PUNCTION: SOMATOSTATIN INHIBITS THE RELEASE OF SOMATOTROPIN.

-- FORCTION: SOCRETON: SECRETED.

-- FORCTION: SOCRETON: SECRETED.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Oncorhynchus mykiss (Rainbow trout) (Salmo gairdneri).
Bukaryota: Metazoa; Chordata; Craniata: Vertebrata; Euteleostomi;
Actinopterygii; Neopterygii: Teleostei; Buteleostei;
Protacanthopterygii: Salmoniformes; Salmonidae; Oncorhynchus.
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  Score 138; DB 1; Length 125;
Pred. No. 6.5e-14;
1; Mismatches 1; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      [TYR7, GLY10] SOMATOSTATIN-28. [TYR7, GLY10] SOMATOSTATIN-14.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 134; DB 1; Length 73
Pred. No. 1.5e-13;
1; Mismatches 0; Indels
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                                                                                            101 STNNLPPRERKAGCKNFYWKGFTSC 125
                                                                   1 SVDNLPPRERKAGCKNFYWKGFTSC 25
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          91.28;
Query Match 93.9%;
Best Local Similarity 92.0%;
Matches 23; Conservative
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Best Local Similarity
Matches 22; Conserv
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Q91194;
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P21780:
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PEPTIDE
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SEQUENCE
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This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way
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leavage on pair of basic residues; Hormone; Signal; Multigene family.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYR21, GLY24 SOMATOSTATIN-28 (POTENTIAL).
TYR7, GLY10 SOMATOSTATIN-14.
"Isolation and characterization of a CDNA encoding for preprosomatostatin containing [Tyr7, Gly10] somatostatin-14 from the endocrine pancreas of rainbow trout, Oncorhynchus mykiss."; Gen. Comp. Endocrinol. 98:253-261(1995).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PUSEC. 1989 (Rel. 10, Created) 01-MAR-1989 (Rel. 10, Created) 01-MAR-1981 (Rel. 17, Last sequence update) 16-OCT-2001 (Rel. 40, Last annotation update) Somatostatin II precursor [Contains: [Tyr21,Gly24]somatostatin-28;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Johnstolatin in pretation (Tragments).
[Ty77,01910] sometostatin-14] (Fragments).
[Ty77,01910] sometostatin-14] (Fragments).
Wyoxocoephalus scorpius (Shorthorn Sculphi) (Daddy sculphi).
Eukaryota; Metazoa; Ohordata; Craniata; Vertebrata; Buteleostomi;
Actinopterygii; Neopterygii; Teleostei; Euteleostei; Neoteleostei;
Acanthomorpha; Acanthopterygii; Percomorpha; Scorpaeniformes;
cottoidei; Cottidee; Myoxocoephaius.
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- FUNCTION: SOMATOSTATIN INHIBITS THE RELEASE OF SOMATOFROPIN.

- PUNCTION: SOMATOSTATIN SECRET.

- SUBCELLULAR LOCATION: Secreted.

- SIMILARITY: BELONGS TO THE SOMATOSTATIN FAMILY.
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"Structural characterization of peptides derived from
prosomatostatins I and II isolated from the pancreatic islets or
species of teleostean fish: the daddy sculpin and the flounder.
Eur. J. Blochem. 168:647-652(1987).
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WIDLINE-87190954; PubMed-2883025;
Cutfield S.M., Carne A., Cutfield J.F.;
"The amino-acid sequences of sculpin islet somatostatin-28 and peptide YY.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 134; DB 1; Length 115;
Pred. No. 2.4e-13;
1; Mismatches 0; Indels
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520595025FCA6D91 CRC64;
                                                                                                                                     I - FUNCTION: SOMATOSTATIN INHIBITS THE RELEASE OF: SUBCELLULIAR LOCATION: SECRETEd.
- SIMILARITY: BELONGS TO THE SOMATOSTATIN FAMILY.
                                                                                                                                                                                                                                                                                                                                                                            use by non-profit institutions as long a
modified and this statement is not removed.
entities requires a license agreement (See No
or send an email to license@isb-sib.ch).
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95.7%;
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104
115 AA;
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1- SUBCELLULAR LOCATION: Secreted.
                                                                          SEQUENCE FROM N.A.
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Matches
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"Characterization of the pancreatic hormones from the Brockmann body
of the tilapla: implications for islet xenograft studies.";
Comp. Blochem. Physiol. 111C:33-44(1995).

- FONCTION: SOMATOSTAIN INTERITS THE RELEASE OF SOMATOTROPIN.

-1- SUMILARITY: BELOWS TO THE SOMATOSTATIN FAMILY.

InterPro: IPRO04250; Somatostatin.

PF03002; Somatostatin. 1.
                                                                                                                                                                                                                                                                                                                                                                          01-NOV-1997 (Rel. 35, Created)
01-NOV-1997 (Rel. 35, Last sequence update)
16-CCT-2001 (Rel. 40, Last annotation update)
16-CCT-2001 (Rel. 40, Last annotation update)
Somatostatin II precursor (Contains: [Tyr21,Gly24]somatostatin-28;
[Tyr7,Gly10]somatostatin-14] (Frapament)
Oreconfromis niloticus (Nile tilapia) (Tilapia nilotica)
Refrayota: Metazoa; Chordate; Craniata; Vertebrata; Euteleostomi;
Actinopterygii; Neopterygii; Teleostei; Buteleostei; Neoteleostei;
Actinopterygiis Actinopterygii; Percomorpha; Perciformes; Labroidei;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SMS2_CARAU STANDARD; PRT; 120 AA.

G9YGH, Q9PUG1,

16-OCT-2001 (Rel. 40, Last sequence update)

16-OCT-2001 (Rel. 40, Last annotation update)

Somatostatin II precursor [Contains: [Tyrz1,G1y24]somatostatin-28;

[Tyr7,G1y10]somatostatin-14].

Garassius auratus (Goldfish).

Carassius auratus (Goldfish).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Carassius auratus (Goldfish).
Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
                 basic residues; Hormone; Multigene family.
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Pred. No. 5e-12;
3; Mismatches 1; Indels
                                                                                 [TYR21, GLY24]SOMATOSTATIN-28.
[TYR7, GLY10]SOMATOSTATIN-14.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                [TYR21, GLY24]SOMATOSTATIN-28. [TYR7, GLY10]SOMATOSTATIN-14.
                                                                                                                                                              11arity 82.1%; Score 123.5; DB 1; Length 12 82.1%; Pred. No. 5.8e-12; Conservative 1; Mismatches 1; Indels
                                                                                                                                6864A59A3FA72C47 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   47C049F4866EF4AC CRC64;
                                                                                                                                                                                                                                                                                                                                                     28 AA.
                                                                                                                                                                                                                                                   1 SVD---NLPPRERKAGCKNFYWKGFTSC 25
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                                                                                                                                                                                                                                                                                                                                                     PRT;
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28 AA; 3155 MW;
                                                                                                                                8036 MW;
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Local Similarity 82.6%;
les 19; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sichlidae, Oreochromis
NCBL_TaxID-8128;
                 pair of
                                                                                                                                                                             Local Similarity es 23; Conserv
PIR; S00166;
Cleavage on
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P81029;
                                                                              PEPTIDE
PEPTIDE
DISULFID
SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Cleavage on pair of basic residues; Hormone; Signal; Multigene family.
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16-0CT-2001 (Rel. 40, Last annotation update)
Somatostatin I precursor [Contains: Somatostatin 26; Somatostatin-14]
Actinopteryg11; Neopteryg11; Teleoste1; Ostarlophys1; Cypriniformes; Cyprinidae; Carassius.
NCBL_TaxID=7957;
                                                                                                                                                                    Lin X.-W., Peter R.E.;
S.Choing and characterization of CDNAs encoding preprosomatostatin-I
and -II from goldfish brain.";
Submitted (JUN-1996) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                    Otto C.J., Peter R.E.;
"The expression of SRIF mRNA in the brain of goldfish.";
submitted (SEP-1997) to the EMBL/GenBank/DDBJ databases.
-I. FUNCTION: SOMATOSTATIN INHIBITS THE RELEASE OF SOMATOTROPIN.
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Bukāryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Actinopterygii; Neopterygii; Amilformes; Amiidae; Amia.
NCBI_raxiD=7924;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             POTENTIAL.
[TYR2], GLY24]SOMATOSTATIN-28.
[TYR7, GLY10]SOMATOSTATIN-14.
BY SIMILARITY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Q -> RW (IN REF. 2).
98957D68011A651A CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                     - FUNCTION: SOMATGSTATIN INTIBITS THE RELEASE OF SOME SUBSTANTIAL SUBSTANTIAL SOME SOFTER SUBSTANTIN FAMILY.
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16-OCT-2001 (Rel. 40, Last anno
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120 AA; 13723 MW;
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5 LPPRERKAGCKNFYWKGFTSC 25
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Q9YGH5;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Hadraws P.C.; Dixon J.E.;
Andrews P.C.; Dixon J.E.;
"Isolation and structure of a peptide hormone predicted from a mRNA sequence. A second somatostatin from the catfish pancreas.";
T Riol Chem. 256:8267-8270(1981).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Bukaryota: Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;
Actinopterygii, Neopterygii; Teleostel; Ostariophysi; Siluriformes;
Ictaluridae; Ictaluria
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            JOINNE-82082515; Pubmed-6171821; Aylor W.L., Collier K.J., Deschenes R.J., Weith H.L., Dixon J.E.; Sequence analysis of a cDNA coding for a pancreatic precursor to
  1- SIMILARITY: BELONGS TO THE SOMATOSTATIN FAMILY.
INTERPO: IPROGASTO. SOMATOSTATIN.
Pfam: PP03002; Somatostatin. 1.
Clarvage on pair of basic residues; Hormone; Multigene family.
NON.TER.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Contains Somatostatin-14 (SS-14)].
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long as its content
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1- SIMILARITY: BELONGS TO THE SOMATOSTATIN FAMILY:
SIMILARITY: BELONGS TO THE SOMATOSTATIN FAMILY. PPro; IPR004250; Somatostatin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                oc. Natl. Acad. Sci. U.S.A. 78:6694-6698(1981).
                                                                                                                                                                                                          Score 107; DB 1,
Pred. No. 6e-10;
1; Mismatches
                                                                                                       SOMATOSTATIN-26.
                                                                                                                                                                                                                                                                                                                                                                                                                                               114 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (Rel. 15, Last sequence update)
(Rel. 15, Last sequence update)
(Rel. 40, Last annotation update
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Somatostatins of the channel catfish.";
dv. Exp. Med. Biol. 188:19-29(1985).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                omatostatin-14 messenger RNA.";
[. Biol. Chem. 257:10372-10377(1982).
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                                                                                                                                                                                                                                                                                                                       6 LAPRERKAGCKNFFWKTFTSC 26
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MEDLINE-85303576; Pubmed-2863931;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MEDLINE-81264223; PubMed-6114953;
                                                                                                                                                                     2931 .MW;
                                                                                                                                                                                                       th 72.8%;
| Similarity 85.7%;
| Similarity 85.7%;
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                                                                                                                                                                     26 AA;
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P01171:
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Actinopterygii; Neopterygii; Teleostei; Ostariophysi; Cypriniformes;
Cyprinidae; Carassius
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16-OCT-2001 (Rel. 40, Last annotation update)
Somatostatin 14 precursor [Contains: Somatostatin-26; Somatostatin-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (NOV-1995) to the EMBL/GenBank/DDBJ databases.
ON: SOMATOSTATIN INHIBITS THE RELEASE OF SOMATOTROPIN.
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SOMATOSTATIN-14.
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B5920015E2D272A4 CRC64;
                                                                                                               of basic residues; Hormone; Signal
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                                                                                                                                                                                                                         62 62 E -> Q (IN REF. 2).
114 AA; 12419 MW; FEE0F2C76F74D99F
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Pred. No. 2.8e-09;
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85.7%;
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A23878.1;
                                                                                                                                                                                                                                                                                                                     Local Similarity 85.7
les 18; Conservative
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es 18; Conserv
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                                                                                                                                              Gallus gallus (Chicken).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostom1;
Archosauria; Aves; Neognathae; Galliformes; Phasianidae; Phasianinae;
               01-OCT-1993 (Rel. 27, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
Somatostatin precursor [Contains: Somatostatin-18; Somatostatin-14].
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Lophius americanus (American goosefish) (Anglerfish).
Bukaryota: Metacoa; Chordata; Cranatata; Vertebrata; Euteleostomi; Actinopterygii; Neoperygii; Tenatata; Euteleostei; Neoteleostei; Acanthomorpha; Paracanthopterygii; Lophiiformes; Lophiidae; Lophius
                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE FROM N.A. PubMed-6107860; Hebrines 1052423; PubMed-6107860; Hobbrt P.M., Crawford R., Shen L., Pictet R., Rutter W.J.; Hobbrt P.M., crawford R., Shen of cDNAs encoding two distinct "Cloning and sequence analysis of cDNAs encoding two distinct somatostatin precursors found in the endocrine pancreas of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 116;
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MEDLINE-81077276; PubMed-6108560;
Goodman R.H., Jacobs J.W., Chin W.W., Lund P.K., Dee P.C.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2; Indels
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21-JUL-1986 (Rel. 01, Created)
21-JUL-1986 (Rel. 01, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
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IGNAL 1 24 BY SIMILARITY.
ROPEP 25 88 BY SIMILARITY.
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1- SIMILARITY: BELONGS TO THE SOMATOSTATIN FAMILY.
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85.7%; Pred. No. 2.9e-09;
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Pfam; PF03002; Somato
                                                                                                                                                                                                                                                                                                                                                        SEQUENCE FROM N.A.
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P01169;
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SMS1_LOPAM
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     #EDLINE-93038702; PubMed-1358069; /audry H., Chartrel N., Conlon J.M.; /audry H., Chartrel N., Conlon J.M.; /aidition of [Pro2,Metl3]somatostatin-14 and somatostatin-14 from the frog brain reveals the existence of a somatostatin gene family in a
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Pfam; PF03002; Somatostatin; 1.
Cleavage on pair of basic residues; Hormone; Multigene family; Signal.

BY SIMILARITY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Enkaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostoml;
Amphila; Batrachia; Anura; Neobatrachia; Ranoidea; Ranidae; Rana
(KSI_TAXID=8406;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TOSTIVINE H., Libramann I., Bucharles C., Vieau D., Coulouarn Y., Fournier A., Conlon J.M., Vaudry H.;
"Occurrence of two somatostatin variants in the frog brain:
"Characterization of the cDNAs, distribution of the mRNAs, and receptor-binding affinities of the peptides.";
Proc. Natl. Acad. Sci. U.S.A. 93:12605-12610(1996).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   chem. Biophys. Res. Commun. 188:477-482(1992). PUCTION: SOMATOSTAIN INHIBITS THE RELEASE OF SOMATOSTROPIN. SUBCELLULAR LOCATION: SOCIETE SOCIETION SOCIETION SUBCELLULAR SECOND TO THE SOMATOSTATIN FAMILY.
                                                                                                                                                                                                                                                                                                16-OCT-2001 (Rel. 40, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
Somatostatin 1 precursor (PSS1) [Contains: Somatostatin-14 (S-I)
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349756FEB4ABE213 CRC64;
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BY SIMILARITY.
SOMATOSTATIN-14.
BY SIMILARITY.
                                                                                                                                                                                                             115 AA.
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LAPRERKAGCKNFFWKTFTSC 114
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            IISSUE-Brain;
MEDLINE-97057290; PubMed-8901629;
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                                                                                                                                                              | LEANRI | Signary | System | 
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P33094;
01-OCT-1993 (Rel. 27, Created)
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MEDLINE-86030691; PubMed-2865169;
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Endocrinology 105:1440-1415(1979).

I FUNCTION: SOMPTOSTATIN INHIBITS THE RELEASE OF SOMATOTROPIN.

SUBJECTIVILAR LOCATION: SCENETION

I SUBJECT S
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Last annotation update)
[Contains: Somatostatin-28; Somatostatin-14]
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Mammalia; Butheria; Cetartiodactyla; Suina; Suidae; Sus.
NCBI_TRAID-9823;
                                                                                                                                                                                                                                                     MEDLINE-80046482; PubMed-387385; Nobe B.D., Spless J., Rivier J.E., Vale W.; **Isolation and characterization of somatostatin from anglerfish panoreatic isjet.**
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              Nucleotide sequence of a cloned structural gene coding for a precursor of pancreatic somatostatin."; vcc. Natl. Acad. Sci. U.S.A. 77:5869-5873(1980).
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"Prosomatostatin 1-64 is a major product of somatostatin gene
expression in pancreas and gut.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 107; DB 1; Length 121;
Pred. No. 3e-09;
1; Mismatches 2; Indels
                                                                                                                                         Goodman R.H., Jacobs J.W., Chin W.W., Lund P.K., Dee P.C. Habener J.F.,
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G -> E (IN REF. 2).
D70C53DC798C2095 CRC64;
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Fam; PF03002; Somatostatin; 1.
:leavage on pair of basic residues; Hormone; Signal;
                                                                                                                                                                                      roc. Natl. Acad. Sci. U.S.A. 79:1682-1682(1982).
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01-CCT-1989 (Rel. 12, Last sequ
16-CCT-2013 (Rel. 40, Last anno
Somatostatin precursor (Contain
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ilarity 85.7%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Schally A.V., Huang W.-Y., Chang R.C.C., Arimura A., Redding T.W., Milar R.P., Hunkapiller M.W., Hood L.E.;
Isolation and structure of pro-somatostatin: a putative somatostatin precursor from pig hypothalamus.",
Proc. Natl. Acad. Sci. U.S.A. 77:4489-4493(1980).
Schmidt W.E., Mutt V., Kratzin H., Carlquist M., Conlon J.M.,
Creutzfeldt W.;
"Isolation and characterization of proSS1-32, a peptide derived from
the N-terminal region of porcine preprosomatostatin.";
FEBS Lett. 192:141-146(1985).
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Schally A.V., Dupont A., Arimura A., Redding T.W., Nishi N.,
Linthkuum G.L., Schlesinger D.H.;
Tisolation and structure of somatostatin from porcine hypothalami.";
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PUGITION: SOMANOSTATIN INHIBITS THE RELEASE OF SOMATOTROPIN.
SUBCELLULAR LOCATION: Secreted.
SIMILARITY: BELONGS TO THE SOMATOSTATIN FAMILY.
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t terminally extended somatostatin: the primary structure of
somatostatin-28.
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fam; PF03002; Somatostatin; 1.
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MEDLINE=80113258; PubMed=7353633;
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dEDLINE-81054799; PubMed-6107906;
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'; PubMed-8988514; Delvalle J., Todisco A., Gantz I., Tong L.,

(See http://www.isb-sib.ch/announce

Somatostatin.

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2; Indels

2; Mismatches

Score 105; DB 1; Length 116; Pred. No. 5.8e-09;

AB49BD231E731C9E CRC64;

SOMATOSTATIN-14 BY SIMILARITY.

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                                                                   Finniss S., vanada T.; 'Contine prosonation of a CDNA, regulation of gene 
"Contine prosonatostatin: isolation of a cDNA, regulation of gene 
expression, and characterization of post-translational processing
                                                                                                                                                                      SOMATOSTATIN INHIBITS THE RELEASE OF SOMATOTROPIN.
                                                                                                                                                                                        -i- SUBCELLULAR LOCATION: Secreted.
-i- SIMILARITY: BELONGS TO THE SOMATOSTATIN FAMILY.
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or send an email to license@lsb-sib.ch)
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Job time : 10.6154 secs
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01-AUG-1992 (Rel. 23, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
Somatostatin precursor [Contains: Somatostatin-28; Somatostatin-14].
                                                                                                                                                                                                                                            MEDLINE-88288237; Pubmed-2899837;
Su C.J., White J.W., Li W.H., Luo C.C., Frazier M.L., Saunders G.F.,
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16-OCT-2001 (Rel. 40, Last annotation update)
Somatostatin precursor [Contains: Somatostatin-28; Somatostatin-14].
                                                                                   cukaryota: Metazoa: Chordata; Craniata; Vertebrata; Buteleostomi;
Mammalia; Butheria; Cetartiodactyla; Ruminantia; Pecora; Boyoidea;
WCBL_TaxID=9913;
                                                                                                                                                                                                                                                                                                                                                                                                   STRAIN-HOISTEIN;
MEDINE-99198780; Pubmed-10100681;
Furu L.M., Razmer G.W., Strausbaugh L., Zinn S.A.;
"Cloning and characterization of the bovine somatostatin gene:";
J. Anim. Sci. 77:492-493(1999).
-I. FUNCTION: SOMATOSTATIN INHIBITS THE RELEASE OF SOMATOTROPIN.
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Mammalia; Eutheria; Carnivora; Fissipedia; Canidae; Canis
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Pred. No. 5.8e-09;
2; Mismatches 2; Indels
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1- SIMILARITY: BELONGS TO THE SOMATOSTATIN FAMILY.
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                                                                                                                                                                                                                                                                                                                Structure and evolution of somatostatin genes.";
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GenCore version 5.1.6 Copyright (c) 1993 - 2003 Compugen Ltd. OM protein - protein search, using sw model Run on: August 13, 2003, 14:48:42; Search time 37:8205 Seconds (without alignments) 100:577 Million cell updates/sec		ber of hits satisfy: B seq length: 0 B seq length: 200000 essing: Minimum Matc	Database : SPTREMBL_23:* 1: sp_archea:* 2: sp_barteria:* 3: sp_fungl:* 4: sp_luman:* 5: sp_lnvertebrate:* 6: sp_namman:* 8: sp_nreman:* 8: sp_organelle:*	+	e greater than or equal to the score of the distriction of the total score distriction of the total score distriction of the total score distriction of the score districti	1 135 91.8 115 13 090743 090943 osteoglossu 2 130 88.4 114 13 090742 090442 patodoon bu 3 123 83.7 28 13 09P0740 09P0740 09P0740 09P0740 0PD0741 0PD0744 0PD0741 0PD0741 0PD0741 0PD0741 0PD0741 0PD0744 0PD0744 <t< td=""></t<>

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Anguilla japonica (Japanese eel),
bukaryotta, Metazoa, Chordata, Craniata; Vertebrata, Euteleostomi,
Actinopterygii, Neopterygii, Teleostel, Anguilliformes; Anguillidae,
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Actinopterygii; Neopterygii; Teleostei; Osteoglossomorpha;
Osteoglossiformes; Notopteridae; Chitala
NOSI_TAXID-112163;
                                                                                                                                                                                                                                                                                                          81.0%; Score 119; DB 13; Length 114; 95.2%; Pred. No. 3.3e-11; arive 0; Mismatches 1; Indels 0
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                                                                                                 Al-Mahrouki A.A., Irwin D.M., Youson J.H.; Romas from several "Characterization of variant somatostatin cDNAs from several osteoglossomorphs: molecular identification and comparative
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     Actinopterygii; Neopterygii; Teleostei; Osteoglossomorpha;
Osteoglossiformes; Mormyridae; Gnathonemus.
                                                                                                                                                                                       Submitted (AUG-2000) to the EMBL/GenBank/DDBJ databases.
EMBL, AF292652; AAK97069.1; -
Interpro; IPR044550; Somatostatin.
Priam; PP0302; Somatostatin; 1.
SEQUENCE 114 AA; 12494 MW; 454DA57A309CA8F2 CRC64;
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AF292653; AAK97070.1;
Pro; IPR004250; Somatostatin.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             InterPro; IPR004250; Somatostati
Pfam; PF03002; Somatostatin; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Somatostatin-related peptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (TrEMBLrel. 19,
(TrEMBLrel. 19,
(TrEMBLrel. 21,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (TrEMBLrel, 13, (TrEMBLrel, 13, TremBlrel, 19,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRELIMINARY:
                                                                                                                                                                                                                                                                                                                                 Local Similarity 95.2 tes 20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRELIMINARY;
                                                                                     SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE FROM N.A.
                                             NCBI_TaxID-42645;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Anguilla.
NCBI_TaxID=7937;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       01-DEC-2001
01-JUN-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                -DEC-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 -MAY-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Submitted
                                                                                                                                                                                                                                                                                                                    Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE
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SEQUENCE.
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Matches
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     SORRETTER
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01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
6SS-28-SOMATOSTAIN-118. 19, Last annotation update)
GSS-38-SOMATOSTAIN-118 peptide
GSS-38-SOMATOSTAIN-118 peptide
Exassius auratus (Goldfish)
Bukaryota: Metazoa: Chordata: Craniata; Vertebrata; Euteleostomi;
Actinopterygii: Neopterygii; Teleostei; Ostarlophysi; Cypriniformes;
Cyprinidae; Carassius
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDLINE-96051491; PubMed-8536941; Gesaka T., Yano K., Yanasaki M., Ando M.; Somatostatin., vasocative intestinal peptide-, and granulin-like peptides isolated from intestinal extracts of goldfish, Carassius
                                                  Pantodon buchholtzi (Butterflyfish),
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Actinopterygil; Neopterygil; Teleostei; Osteoglossomorpha;
Osteoglossiformes; Pantodontidae; Pantodon.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      090741 PRELIMINARY; PRT; 114 AA.
090741, 01-DEC-2001 (TrEMBLE1. 19, Created)
01-DEC-2001 (TrEMBLE1. 19, Last sequence update)
01-MAR-2002 (TrEMBLE1. 20, Last annotation update)
Preprosmatostatin.
Gnathonemus petersii.
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
                                                                                                                                                                                                                                                                                                                                                                                                                 Match 88.4%; Score 130; DB 13; Length 114; Local Similarity 95.5%; Pred. No. 5.7e-13; Mismatches 0; Indels 0
                                                                                                                                                                                                   Al-Mahrouki A.A., Irwin D.M., Youson J.H.; Characterization of variant somatostatin cDNAs from several osteoglossomorphs: molecular identification and comparative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 123; DB 13; Length 28; Pred. No. 1.8e-12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2; Indels
                                                                                                                                                                                                                                                                                   Submitted (AUG-2000) to the EMBL/GenBank/DDBJ databases EMBL; AF292651; AAK97068.1; -InterPro: IPR004250; Somatostatin.
                                                                                                                                                                                                                                                                                                                                                       102; Somatostatin; 1.
114 Aa; 12352 MW; 7E3D44CB6A27B12F CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            15D271F677C945BE CRC64;
Last sequence update)
Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             28 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gen. Comp. Endocrinol. 99:298-306(1995).
InterPro; IPR004250; Somatostatin.
Pfam; PF03002; Somatostatin; 1.
SEQUENCE 28 AA; 3204 MW; 15D271F677cs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   4 SSNHLPARERKAGCKNFYWKGFTSC 28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 SVDNLPPRERKAGCKNFYWKGFTSC 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               93 NVPPRERKAGCKNFYWKGFTSC 114
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          · PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           4 NLPPRERKAGCKNFYWKGFTSC 25
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19,
20,
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21; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRELIMINARY;
(TrEMBLrel.
                                          Preprosomatostatin
                                                                                                                                                                                  SEQUENCE FROM N.A.
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                                                                                                                                                                                                                                                                                                                                                Pfam; PF03002;
SEQUENCE 114
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Matches
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Gaps

Indels

Length 120;

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Pfam; PF03002; Somatostatin; 1
SEQUENCE 120 AA; 13783 MW; 00828D35263E8805 CRC64;
                                                                                                                                        Match 72.8%; Score 107; DB 13; Local Similarity 72.0%; Pred. No. 2.9e-09; les 18; Conservative 3; Mismatches 4;
                                                                                                                                                                                                                                                                              1 SVDNLPPRERKAGCKNFYWKGFTSC 25
2654; AAK97071.1; -.
IPR004250; Somatostatin.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NCBI_TaxID-7904;
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                                                                                                                                                Ouery Match
Best Local S
Matches 18
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Brachydanio rerio (Zebrafish) (Danio rerio).
Eukaryota: Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Actinopterygil; Neopterygil; Teleostel; Ostariophysi; Cypriniformes;
Cyprinidae; Danio.
NCBI_TaxiD-7955;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Catostomus commersoni (White sucker).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Actinopterygii; Neopterygii; Teleostei; Ostariophysi; Cypriniformes;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             expression of two somatostatin genes during zebrafish
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  -22045842; PubMed-12049777; ... Deflorian G., Biemar F., Bortolussi M., Martial J.A.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        72.8%; Score 107; DB 13; Length 114; 85.7%; Pred. No. 2.7e-09; 1ve 1; Mismatches 2; Indels 0
                                                                                                                                                                                                                                                                  77.6%; Score 114; DB 13; Length 25; 84.0%; Pred. No. 4.3e-11;
                                                                                                                                                                                                                                                                                                                                       4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Al-Mahrouki A.A., Irwin D.M., Youson J.H.,
"Molecular Coining and characterization of white sucker
preprosomatostatin ",
Submitted (AUG-2000) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       InterPro; IPR004250; Somatostatin.
Pfam: PF03002; Somatostatin; 1.
SEQUENCE 114 AA; 12416 MW; 3D41424AE54E7408 CRC64;
                                                                ion and water absorption across the intestine of the L. Exp. Biol. 188:205-216(1994).
InterPro: IPR004250; Somatostatin.
Pfam: PF003002; Somatostatin; 1.
SEQUENCE 25 AA; 2860 MW; BFC672143A04A3F5 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              01-CCT-2002 (TrEMBLrel. 22, Created)
01-CCT-2002 (TrEMBLrel. 22, Last Sequence update)
01-MAR-2003 (TrEMBLrel. 23, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           01-DEC-2001 (TIEMBLrel. 19, Created)
01-DEC-2001 (TIEMBLrel. 19, Last sequence update)
01-MAR-2002 (TIEMBLrel. 20, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            114 AA.
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                                                                                                                                                                                                                                                                                 84.0%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                  1 SVDNQQGRERKAGCKNFYWKGPTSC 25
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EMBL; AF435965; AAM54072.1; --
EMPL; AF43596; AAM54072.1; --
EMPL; AF43596; AAM54072.1; --
EMPL; AAM54072.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                5 LPPRERKAGCKNFYWKGFTSC 25
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Best Local Similarity 85.79
watches. 18; Conservative
                                                                                                                                                                                                                                                                                                 Local Similarity 84.0 ses 21; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRELIMINARY;
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   Gesaka T., Yano K., Ya
"Somatostatin-related
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE FROM N.A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Somatostatin-14.
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                                                                                                                                                                                                                                                                  Query Match
Best Local S
Matches 21
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Q90x39;
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08JHX5
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Q90X39

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Trabucchi M., Tostivint H., Lihrmann I., Sollars C., Vallarino M., Dores K.M., Yaudry H.; Polygenic expression of somatostatin in the sturgeon Acipenser transmontanus: molecular cloning and distribution of the mRNAs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                M., Tostivint H., Lihrmann I., Sollars C., Vallarino M.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Actinopterygil; Chondrostei; Acipenseriformes; Acipenseridae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Dores R.M., Vaudry H.;
"Polygenic expression of somatostatin in the sturgeon Acipenser
"Polygenic expression of somatostatin in the sturgeon Acipenser
transmontanus: molecular cloning and distribution of the mRNAs
encoding two somatostatin precursors.";
Comp. Neurol. 0:0-0(2001).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ö
                                                                                                                                                                                                                                                                                                                  Acipenser transmontanus (White sturgeon).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleost
Actinopterygil; Chondrostei; Acipenseriformes; Acipenseridae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 116;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               F03002; Somatostatin; 1.
E 116 AA; 12616 MW; 72E0C3FF6C80650F CRC64;
                                                                                                                                                                              090XEI;
01-DEC-2001 (TYEMBLIZEL 19, Created)
01-DEC-2001 (TYEMBLIZEL 19, LAST SEQUENCE UPDATE)
01-MAR-2002 (TYEMBLIZEL 20, LAST annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     01-DEC-2001 (TrEMBLrel: 19, Created)
01-DEC-2001 (TrEMBLrel: 19, Last sequence update)
01-MAR-2002 (TrEMBLrel: 20, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            71.4%; Score 105; DB 13;
81.0%; Pred. No. 5.8e-09;
11ve 2; Mismatches 2;
                                                                                                                                                 116 AA.
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96 NINOLYPRERKAGCKNFFWKIFISC 120
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                                                                                                                                                 PRT;
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                                                                                                                                                          PRELIMINARY;
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Acipenser transmontanus
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Gaps

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Eukaryota; Matazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
NCBL_raxID=9606;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalla, Eutheria, Primates, Catarrhini, Hominidae, Homo
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       Pred. No. 0.00034; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Strausberg R.; submitted (NOV-2002) to the EMBL/GenBank/DDBJ databases. EMBL; BC040034, AAH40034.1; - 1 1 1 1 1 1 1 ^{-1}
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)2) to the EMBL/GenBank/DDBJ databases
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164 AA; 19047 MW; 389682C41252426A CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE 122 AA; 13369 MW; A1279CA09CB0CB44 CRC64;
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(TrEMBLrel. 23, Last sequence update)
(TrEMBLrel. 23, Last annotation update)
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Last annotation updat
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Pred. No. 0.015;
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40.7%; Pred. No.
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InterPro; IPR004250; Somatostatin.
Pfam: PP03002; Somatostatin.
Hypothetical protein.
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01-OCT-2002 (TrEMBLrel. 22, Lau
01-MAR-2003 (TrEMBLrel. 23, Lau
Hypothetical protein.
Homo sapiens (Humer
                                                                                              91 ERKTGCKNYFWKSRTAC 107
                                                                   9 ERKAGCKNFYWKGFTSC: 25
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Best Local Similarity 40.7%;
Matches 11; Conservative
       Best Local Similarity 64.7%;
Matches 11; Conservative
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Best Local Similarity 40.7
Matches 11; Conservative
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EMBL, AF521016; AP.
InterPro.
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Q9SGS6
ID Q9SGS6
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08IUV6
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08NFE5
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Eukaryota, Metazoa: Chordata, Craniata; Vertebrata; Euteleostomi;
Actinopterygii, Neopterygii; Teleostei; Ostariophysi; Cypriniformes;
Cyprinidae; Danio.
                                                                                                                                                                                                                                                                                                                                                                                                                                        Anyulla igaponica (Japanese eel).
Ebkaryota: Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi
Actinopterygii; Neopterygii; Teleostei; Anguilliformes; Anguilli
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ., Zecchin E., Bortolussi M.; earance of pancreatic hormone-expressing cells in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 85.5; DB 13; Length 23;
Pred. No. 1.5e-06;
0; Mismatches 4; Indels
                                                                                      Length 111;
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                                                                                         Db _____6, Indels
InterPro; IPR004250; Somatostatin.
Pfam; PF03002; Somatostatin; 1.
SEQUENCE 111 AA; 12748 MW; 4E27DB90896A9025 CRC64;
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rvo; IPR004250; Somatostatin.
PPF03002; Somatostatin; l.
CE 107 AA; 11839 MW; E12C923E56642EFB CRC64;
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O9PRV6;
01-MAY-2000 (TTEMBLrel. 13, Last sequence update)
01-JUN-2002 (TTEMBLrel. 13, Last sequence update)
01-JUN-2002 (TTEMBLrel: 21, Last annotation update)
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Last annotation update)
                                                                                  Score 92; DB 13;
Pred. No. 6.7e-07;
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                                                                               Query Match 62.6%; Score 92; DB Best Local Similarity 66.7%; Pred. No. 6.7e Matches 16; Conservative 2; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MEDIINE-95003944; PubMed-7765422;
UGGSKa T., Yano K., Yamasaki M., P
Zool, Sci. 11:491-494(1994).
NOW_TER
                                                                                                                                                                                                                                                                                                                                                                                                                     (Fragment
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Local Similarity 79.2%;
les 19; Conservative
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Q9DDE4;
Q1-MAR-2001 (TEMBLE-1 16,
Q1-MAR-2001 (TEMBLE-1 16,
01-OCT-2002 (TEMBLE-1 22,
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NCBI_TaxID=7937;
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Pfam; PF03002;
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SEQUENCE

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RESULT 12 Q9DDE4

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SEQUENCE

RESULT 11

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Pr 095656;

Dr 1-MXY-2000 (TrEMBLrel. 13, Last sequence update)

Dr 10-MXY-2000 (TrEMBLrel. 13, Last sequence update)

Sequence Trems (Tremserl. 13, Last sequence update)

Sequence Tremserl. 13, Last sequence update)

Cr Carborous Tremserl. 13, Last sequence update)

Sequence Tremserl. 14, Tradispophyta; Embryophyta; Tracheophyta; Seperatophyta; Magnollophyta; endicotyledons; core endicots; Rosidae; core encretage II; Brasslcales; Brasslcaceae; Arabidopsis. 11

Rederspiel N.A., Palm C.J., Conway A.B., Conn L., Hansen N.F., R. Altefil H., Mayuyen M., Lam B., Buehler E., Dunn P., Gonzalez A., Schwartz J., Shinn P., Toriumi M., Vysotskala V.S., Marker M., Yu G., A. Schwartz J. Shinn P., Toriumi M., Vysotskala V.S., Marker M., Yu G., Exer. J., Theologis A., Davis R.W.;

R. Schwartz J., Theologis A., Davis R.W.;

DR Romartz J., Theologis S. Zaf_FPHD.

DR Romarts SMO029; PRD: Zaf_FPHD.

DR Romarts
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Search completed: August 13, 2003, 14:52:59 Job time: 38.8205 secs

14 3 AAP20165 Somatostatin. 5 14 AAP40366 Sequence of system o	77 86.5 14 19 77 86.5 14 19 77 86.5 14 20 77 86.5 14 20 77 86.5 14 20 77 86.5 14 20 77 86.5 14 21 77 86.5 14 21 86.5 14 21	6.5 14 21 AAY66596 Peptide sequent AAV66596 Peptide sequent AAV66596 Peptide sequent AAV66596 Peptide sequent AAV66745 Peptide sequent AAV67145 Pe	77 86.5 14 22 AAB45659 ALIGNMENTS 07665 standard; Peptide; 14 AA.	AAU07665; 04-DBC-2001 (first entry) Rainbow trout somatostatin 14 (SS-14) variant peptide sequence. Rainbow trout; somatostatin; preprosomatostatin; hypersecretion; PPSS-I; PPSS-II'; endocrine tumour; pituitary gland; glucagonoma; AIDS; gastroenteropancreatic tissue; acromegaly; gastrinoma; diabetes mellitus;	carcinoid syndrome; cell proliferation; apoptosis; growth hormone; SS-14; glucagon; acquired immunodefluciency syndrome; neurological disorder; HIV; epilepsy; Alzheiner's disease; Huntington's disease; neuroprotective; neoplasm; metastasis; gene therapy; antidiabetic; nootropic; cytostatic; anti-human immunodeficiency virus; osteopathic; anticonvulsant. Oncorhypchus mykiss.	03-JUN-2001. 01-DEC-2000; 2000CA-2325169. 03-DEC-1999; 99US-0168934. (NDSU-) NDSU RES FOUND. Sheridan MA, Moore CA, Kittelson JD;
			45 RESULT 1 AAU07665 ID AAU XX	K K K K K K K K K K K K K K K K K K K	KW KW KW XX XX PN	XX X X X X X X X X X X X X X X X X X X
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GenCore version 5.1.6 Copyright (c) 1993 - 2003 Compugen Ltd. OM protein - protein search, using sw model Run on: August 13, 2003, 14:45:37; Search time 26.2051 Seconds (without alignments) 84.799 Million cell updates/sec Title: Berfect score: 89 Sequence: 1 AGCKNFYWKGFTSC 14 Scoring table: BLOSUM62	Gapop 10.0 , Ga 1107863 seqs, Juber of hits satisfying DB seq length: 0 DB seq length: 2000000	TX S	nneseqp-embl/AA1985. Ineseqp-embl/AA1986. Ineseqp-embl/AA1980. eneseqp-embl/AA1989. eneseqp-embl/AA1999. eneseqp-embl/AA1991. eneseqp-embl/AA1993. eneseqp-embl/AA1993.	10. SIDSI/qcgdata/qeneseq/qeneseqp-embl/AA1996.DAT:* 17. /SIDSI/qcgdata/qeneseqq-emeseqp-embl/AA1996.DAT:* 18. /SIDSI/qcgdata/qeneseq-qeneseqp-embl/AA1997.DAT:* 19. /SIDSI/qcgdata/qeneseq-qeneseqp-embl/AA1999.DAT:* 20. /SIDSI/qcgdata/qeneseq-qeneseqp-embl/AA1999.DAT:* 21. /SIDSI/qcgdata/qeneseq-qeneseqp-embl/AA2000.DAT:* 22. /SIDSI/qcgdata/qeneseq-qeneseqp-embl/AA2000.DAT:* 23. /SIDSI/qcgdata/qeneseq-qeneseqp-embl/AA2002.DAT:* 24. /SIDSI/qcgdata/qeneseq-qeneseqp-embl/AA2002.DAT:*	Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution. SUMMARIES Result Query No. Score Match Length DB ID Description	1 89 100.0 14 22 AAU07665 Rainbow trout some 2 89 100.0 111 22 AAU07667 Rainbow trout prep 4 89 100.0 115 2 AAU07667 Rainbow trout prep 5 83 93.3 14 7 AAP60190 Sequence of preproduce of

		æ			SUMMAKIES	
Result No.	Score		Ouery Match Length DB	08	ΙD	Description
-	89	100.0	14	22	AAI107665	cano tuori modei ag
	68	100.0		22	AAD07668	Rainbow trout prep
m	68	100.0	115	22	AAU07667	1 -
4	83	100.0	•	m	AAP20029	of or or
5	83	93.3		7	AAP60190	Sometostation 28 an
9	83	93.3	28	7	AAP61714	Sometostatin-28 an
. 1	83	93.3	28	10	AAP90989	Analogue of angler
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σ	77	86.5	14	ď	AAP10421	Somatostatin glyco
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	13	77	86.	i.	14	2	AAP40808		
	14	11	86.	S	14	7	AAP60863		ď
	15	11	86.	2	14	8	AAP70928		Oma
	16	77	86.	2	14	01	AAP94654		
	17	77	86.	5	14	13	AAR30543		151
	18	77	86.	S	14	14	AAR38652		Sequence encoded b
	19	77	.86	D.	14	15	AAR50276		Somatostatin S14.
	20	77	86.	'n	. 14	16	AAW01514		
	21	77	86.	Ŋ	14	16	AAR95412		Somatostatin-14.
		77	86.	2	14	11	AAR92932		Somatostatin pepti
	23	11	86.	ς.	14	19	AAW68299		
		77	86	2	14	19	AAW51858.		Somatostatin analo
		77	86.	ις O	14	13	AAW50946		Somatostatin analo
	56	77	86.	G	14	19	AAW50940		Somatostatin analo
	27	11	86.	Ŋ	14	19	AAW44018		Corticostatin pept
:	28	17		2	14	20	AAY50231		υ
	53	11	86.	ú	14	20	AAX39620		CTLA-4 VLD CDR 100
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	31	11	86.	N	14	50	AAW97182		Somatostatin pepti
	32	11	86.	S	14	71	AAB08301		Amino acid sequenc
	33	77	86.	2	14	21	7		Human somatostatin
	34	11	86.	s S	14	21	AAY91200	,	Somatostatin, SEQ
	35	77	. 98	D.	14	21	AAY68596		Peptide sequence o
	36	77	86.	2	14	22	AAU05145		Receptor-selective
	37	7.7	86.	ro O	14	22	AAB83200		Human somatostatin
	38	77	98		14	.55	AAB91001		
	39	77	86.	5	14	22	AAB91015	•	
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RES	RESULT 1								
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5 X	04 - DEC - 2001	1007-	H)	(rirst	entry)	2			
DE	Rainbo	Rainbow trout		omato	somatostatin	in 14	(SS-14)	variant peptide	sednence.
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The invention relates to an Oncorhynchus mykiss somatostatin polypeptide containing a portion of preprosomatostatin I (PSSS-1) and/or a portion of preprosomatostatin II (PSS-1) and/or a portion of preprosomatostatin II (PSS-II). The protein sequences and their associated polynuclectides are useful for identifying modified somatostatin polypeptide which functions as a somatostatin agonist useful for research, therapeutics or diagnostics, including medical and veterinary applications. The wild-type somatostatin and its modified the pituitary (e.g. acromegaly) or gastroenteropancreatic tissues (e.g. qastroin are useful for treating hy or gastroenteropancreatic tissues (e.g. gastruinam, glucagonoma, carcinold syndrome), to cause tumour shrinkage through their effects on cell proliferation and apoptosis and as adjuncts and all the treatment of diabetes mellitus via an absorbation of growth hormone
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Rainbow trout; somatostatin; preprosomatostatin; hypersecretion; PPSS-I; PPSS-II; PPSS-III; endocrine tumour; pituitary gland; glucagonoma; AIDS; gastroenteropancreatic tissue; acromegaly; gastrinoma; dibbetes mellitus; carcinoid syndrome; cell proliferation; apoptosis; growth hormone; glucagon; acquired immunodeficiancy syndrome; neurological disorder; HIV; epliepsy; Alzheimer's disease; Huntington 's disease; meuroprotective; neoplasm; metastasis; gene therapy; antidiabetic; nootropic; cytostatic; anti-human immunodeficiency virus; osteopathic; anticonvulsant.
                                                                                                                                                                                                                                                                                                                                                                                                                                                 and glucagon. In addition, dysfunctional somatostatin secretion is associated with acquired immunodeficiency syndrome (AIDS) and various neurological disorders (e.g. epilepsy, Alzheimer's disease and Huntington's disease) and somatostatin antagonists are effective in the treatment of such conditions. Nucleic acids encoding the polypeptides are useful in gene therapy and fusion peptides can be targeted to neoplasms and their metastases, inhibiting the release of their secretory products. This sequence represents a variant somatostatin 14 (SS-14) peptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Rainbow trout preprosomatostatin II (PPSS-II'') polypeptide.
                                                             New somatostatin polypeptides derived from Oncorhynchus for treating diabetes mellitus, acromegaly, gastrinoma, Immunodeficiency syndrome and neurological disorders
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      "PPSS-II'' pre-sequence"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        "PPSS-II'' pro-sequence"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     .
111
hte= "Mature PPSS-II''"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Location/Qualifiers
1..25
/note= "Signal peptide"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAU07668 standard, Protein; 111 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ó
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity 100.0%;
Matches 14; Conservative 0;
                                                                                                                                                  Claim 2; Fig 6; 52pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 AGCKNFYWKGFTSC 14
||||||||||||||||||||1
1 AGCKNFYWKGFTSC 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ..97
/note= "
87
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /note=
87..97
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1..86
/note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Oncorhynchus mykiss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       14 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Key
Peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Pept1de
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 2
AAU07668
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containing a portion of prepresentation in the solution of prepresentation in the solution of prepresentation if (PPSS-II) and/or a portion of prepresentation if (PPSS-II) and/or a portion of associated polynucleotides are useful for identifying modified somatostatin polypeptide which functions as a somatostatin agonist useful for research, therapeutics or diagnostics, including medical and veterinary applications. The wild-type somatostatin and its modified version are useful for treating hypersecretion from endocrine tumours in the pituitary (e.g. acromegaly) or gastroenteropancreatic tissues (e.g. qastroenteropancreatic tissues (e.g. qastroin and apoptosis and as adjuncts in the treatment of diabetes mellitus via inhibition of growth hormone
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              somatostatin polypeptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       treatment of such conditions. Nucleic acids encoding the polypeptides are useful in gene therappy and fusion peptides can be targeted to neoplasms and their metastases, inhibiting the release of their secretory products.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PPSS-II'; PPSS-II'; endocrine tumour; pituitary gland; glucagonoma; AIDS;
gastroenteropancreatic tissue; acromegaly; gastrinoma; diabetes mellitus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           in the treatment of diabetes mellitus via inhibition of growth hormone and glucagon. In addition, dysfunctional somatostatin secretion is associated with acquired immunolementy syndrome (AIDS) and various neurological disorders (e.g. epilepsy, Alzheimer's disease and Huntington's disease) and somatostatin antagonists are effective in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Rainbow trout; somatostatin; preprosomatostatin; hypersecretion; PPSS-I; PPSS-II'; endocrine tumour; pituitary gland; glucagonoma; AIDS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   is sequence represents O. Mykiss PPSS-II'' protein.
te: The features for this sequence are specifically claimed in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New somatostatin polypeptides derived from Oncorhynchus mykiss, u
for treating diabetes mellitus, acromegaly, gastrinoma, acquired
Immunodefictency syndrome and neurological disorders -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Rainbow trout preprosomatostatin II (PPSS-II') polypeptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 100.0%; Score 89; DB 22;
100.0%; Pred. No. 4.7e-06;
tive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     to an Oncorhynchus mykiss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        e release of
PPSS-II'' pro
                                                     98.111
/note= "SS-14 variant peptide"
                          "Dibasic cleavage site"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Moore CA, Kittelson JD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  .AAU07667 standard; Protein; 115 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 1; Fig 3; 52pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        98 AGCKNFYWKGFTSC 111
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 ACCKNFYWKGFTSC 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                04-DEC-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity 100.
96..97
                                  /note-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2001-425997/46.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention relates
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        111 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       N-PSDB; AAS12935.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       specification.
Cleavage-site
                                                                                                                                                        CA2325169-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sheridan MA,
                                                                                                                                                                                                                                                                                                                                                                                                           UDSU-) NDSU
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAU07667;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
                                                         Peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 89; DB 22; Length 14;
Pred. No. 6.5e-07;
Mismatches 0; Indels
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Length 111;

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Pred. No. 4.9e-06;

100.08;

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The invention relates to an Oncorhynchus mykiss somatostatin polypeptide containing a portion of preprosomatostatin I (PSS-I) and/or a portion of preprosomatostatin II (PPSS-II). The protein sequences and their associated polynucleotides are useful for identifying modified somatostatin polypeptide which functions as a somatostatin agonist useful for research, therapeutics or diagnostics, including medical and veterinary applications. The wild-type somatostatin and its modified the pituitary (e.g. acromeaply) or gastroenteropancreatic tissues (e.g. question are useful for treating hypersecretion from endocrine tumours in the pituitary (e.g. acromeaply) or gastroenteropancreatic tissues (e.g. gastroin and acrinical syndrome), to cause tumour shrinkage through their effects on cell proliferation and apoptosis and as adjuncts and the preatment of diabetes mailitus via inhibition of growth hormone
                                                                   cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         release of their secretory products
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                and giucagon. In auguitou, uyaiumiciouse associated with acquired immunodeficiency syndrome (AIDS) and various associated with acquired immunodeficiency syndrome (AIDS) and various meurological disorders (e.g. peliepsy, Alzheimer's disease and Huntington's disease) and somatostatin antagonists are effective in the treatment of such conditions. Nucleic acids encoding the polypeptides as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New somatostatin polypeptides derived from Oncorhynchus mykiss, useful for treating diabetes mellitus, acromegaly, gastrinoma, acquired immunodeficiency syndrome and neurological disorders
  carcinoid syndrome; cell proliferation; apoptosis; growth hormone; glucagon; acquired immunodeficiency syndrome; neurological disorder; epilepsy; Alzheimer's disease; Huntington's disease; neuroprotective; neoplasm; metastasis; gene therapy; antidiabetic; nootropic; cytostatianti-human immunodeficiency virus; osteopathic; anticonvulsant.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     is sequence represents O. Mykiss PPSS-II' protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   nt of such conditions. Nucleic acids encoding the point gene therapy and fusion peptides can be targeted in metastases, inhibiting the release of their secre
                                                                                                                                                                                                                                                                                                                                                                                                             /note= "Prosomatostatin II'"
1100..101
/note= "Dibasic cleavage site"
1102..115
/note= "SS-14 variant peptide"
                                                                                                                                                                                                                                           note- "PPSS-II' pre-sequence"
                                                                                                                                                                                                                                                                                                                                              . 101
ote- "PPSS-II' pro-sequence"
                                                                                                                                                                                                                                                                                                                                                                                     26.115
/note= "Mature PPSS-II'"
                                                                                                                                                                                                       'note- "Signal peptide"
                                                                                                                                                                                                                                                                                                                             'note- "Encoded by CAA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Kittelson JD;
                                                                                                                                                                      Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 2; Fig 3; 52pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       01-DEC-2000; 2000CA-2325169.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                99US-0168934
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Moore CA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NDSU-) NDSU RES FOUND.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2001-425997/46.
N-PSDB; AAS12934.
                                                                                                                        Oncorhynchus mykiss.
                                                                                                                                                                                                                                                                                                          Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         and glucagon. Irassociated with
                                                                                                                                                                                                                                                                                                                                                                                                                                Cleavage-site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CA2325169-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sheridan MA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              03-DEC-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              03-JUN-2001.
                                                                                                                                                                      Key
Peptide
                                                                                                                                                                                                                             Protein
                                                                                                                                                                                                                                                                    Protein
                                                                                                                                                                                                                                                                                                                                                   Peptide
                                                                                                                                                                                                                                                                                                                                                                                          Pept1de
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Peptide
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Gaps
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                                ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ;
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The inventors claim preprosomatostatin-1, prosomatostatin-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  beptide which surprisingly contains a 14 AA sequence of a peptide which surprisingly contains a 14 AA sequence at its terminus which differs from 51 by only 2 AAs, and is termed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 125;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New somatostatin and precursors - produced by transformed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Somatostatin; growth hormone; peptide hormone; secretion
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence of preprosomatostatin-2 encoded on pLaS2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (Updated on 16-AUG-2002 to add missing OS field.) (Updated on 25-MAR-2003 to correct PA field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Ouery Match
100.0%; Score 89; DB 3; I
Best Local Similarity 100.0%; Pred. No. 5.3e-06;
Matches 14; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Pictet RL, Rutter WJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Location/Qualifiers
112..125
/label- Somatostatin II
                                                                                                                                                                                                                                                AAP20029 standard; Protein; 125 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example; Fig 3; 50pp; English.
                                                                                                      102 AGCKNFYWKGFTSC 115
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       80US-0181046.
                                                                                                                                                                                                                                                                                                                                                    (updated)
(updated)
(first entry)
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                                                                         1 AGCKNFYWKGFTSC 14
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                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (REGC ) UNIV CALIFORNIA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Hobart P, Crawford R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 1982-18113E/10.
N-PSDB; AAN20034.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Lophius americanus.
Best Local Similarity
Matches 14; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           microorganisms
                                                                                                                                                                                                                                                                                                                                                 25-MAR-2003
16-AUG-2002
14-AUG-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    03-MAR-1982.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     21-AUG-1981;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  atostatin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       25-AUG-1980;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     EP46669-A.
                                                                                                                                                                                                                                                                                                  AAP20029;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Protein
                                                                                                                           g
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Key Modified-site

Synthetic.

25-MAR-2003 28-JUL-1991

AAP60190;

RESULT 5

31-AUG-1984; 01-APR-1987; 16-AUG-1985;

05-MAR-1986. EP173527-A.

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Angler fish sometostatin-28 and analogue and fragment - useful in
inhibiting insulin secretion in insulinoma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Somatostatin 28; SS-28; analogue; insulin secretion inhibitor; insulinoma; gastric acid secretion; thermoregulation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 28;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      somatostatin-14 or somatostain-28, which is more potent the secretion for treatment of insulinoms (Updated on network)
                                                                                                                                                                                                                                                                                                                                                                                                                                                               The protein sequence is an insulin-selective analogue of anglerfish somatostatin-28, which is more potent the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 83; DB 7;
Pred. No. 1e-05;
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (Updated on 25-MAR-2003 to correct PR field.) Updated on 25-MAR-2003 to correct PA field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            note-"Bonded to Cys-28"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 'note-"Bonded to Cys-17"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Analogue of anglerfish somatostatin 28.
                                                                                                                                                                                                                                        (SALK ) SALK INST BIOLOGICAL STUDIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /note="Also claimed"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAP90989 standard; peptide; 28 AA.
Misc-difference 23
/label= Hyl, Lys
                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 4; Page 18; 19pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          label-Lys, Hyl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 93.3%;
                                                                                                                                             85EP-0305867.
                                                                                                                                                                                84US-0646610
87US-0033295
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          87US-0033295
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 AGCKNFYWKGFTSC 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       15 AGCKNFYWXGFTSC 28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (updated)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Ouery Match
Best Local Similarity 92.9
Matches 13, Conservative
                                                                                                                                                                                                                                                                                                                        WPI; 1986-063363/10.
                                                                                                                                                                                                                                                                                   Spiess J, Noe BD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 28 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       disc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Key
Disulfide-bond
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Disulfide-bond
                                                                                                                                        16-AUG-1985;
                                                                                                                                                                                  31-AUG-1984;
31-APR-1987;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   25-MAR-2003
08-JUN-1990
                                                                                                     05-MAR-1986.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          01-APR-1987;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US4816438-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  28-MAR-1989.
                                                             EP173527-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Anglerfish.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAP90989;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ò
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Angler fish somatostatin-28 and analogue and fragment - useful in
inhibiting insulin secretion in insulinoma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 83; DB 7; Length 14;
Pred. No. 5.3e-06;
0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The protein sequence is an insulin-selective analogue of anglerfish somatostatin-28, which is more potent than secretion for treatment of insulinoma. Secretion for treatment of insulinoma. (Updated on 25-MAR-2003 to correct PR field.) (Updated on 25-MAR-2003 to correct PR field.)
                                                                                                                                                                                                                        Somatostatin-28; insulin-selective; insulinoma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Somatostatin-28; insulin-selective; insulinoma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (SALK ) SALK INST BIOLOGICAL STUDIES.
                                                                                                                                                                                                                                                                                                        Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Location/Qualifiers
                                     AAP60190 standard; protein; 14 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAP61714 standard; Protein; 28 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 4; Page 18; 19pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      93.3%;
92.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 85EP-0305867.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       84US-0646610.
87US-0033295.
                                                                                                                                                                                                                                                                                                                                              /label- Hyl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  25-MAR-2003 (updated)
28-JUL-1991 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 AGCKNFYWXGFTSC 14
                                                                                                                   (updated)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 AGCKNFYWKGFTSC 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity 92.9°
Matches 13; Conservative
                                                                                                                                                                              Somatostatin-28 analogue.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Somatostatin-28 analogue.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 1986-063363/10.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Spiess J, Noe BD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 14 AA;
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Synthetic.

Key

AAP61714;

RESULT AAP6171

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O; Gaps

1; Indels

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/label- Ri
/note- "May be opt. substd. by H or a hexose or
aminohexose motety modified in the 2-position
with an amide group, the hexose having the
pyranose structure; RI-R4 are not all H"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /label- R2
/note= "May be opt. substd. by H or a hexose or
aminohexose modety modified in the 2-position
with an amide group, the hexose having the
pyranose structure; R1-R4 are not all H"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /label- R3
/note= "May be opt. substd. by H or a hexose or aminohexose moiety modified in the 2-position with an amide group, the hexose having the pyranose structure; R1-R4 are not all H"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     "May be opt. substd. by H or a hexose or
aninohexose molety modified in the 2-position
with an amide group, the hexose having the
pyranose structure; R1-R4 are not all H*
                             This peptide is an example of a generic formula for somatostatin derive. which inhibit secretion of growth hormone and are useful itreat diabetes mellitus, acromegaly, angiopathy and in diagnosis. See AAP10308-P10348.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Carbohydrate molety; pentose; hexose; pyranose; glucose; fructose
                                                                                                                                                                                                  Score 77; DB 2;
Pred. No. 4.3e-05;
                                                                                           See AAP10308-P10348.
(Updated on 25-MAR-2003 to correct PR field.)
                                                                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Somatostatin glycosylated analogue.
                                                                                                                                                                                                                                                                                                                                                                                                                                    AAP10421 standard; peptide; 14 AA.
                                                                                                                                                                                                  86.5%;
llarity 85.7%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Guillemin RCL, Lavielle S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (SALK-) SALK INST BIOLOGIC.
                                                                                                                                                                                                                                                                                   1 AGCKNFYWKGFTSC 14
                                                                                                                                                                                                                                                                                                            17-DEC-1992 (first entry)
                                                                                                                                                                                                              Local Similarity
ses 12; Conserv
                                                                                                                                                           Sequence 14 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disulphide_bond
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Key
Modified-site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Modified-site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Modified-site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Modified-site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      08-NOV-1979;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                08-NOV-1979;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US4280953-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAP10421;
                                                                                                                                                                                                    Query Match
                                                                                                                                                                                                                           Best Loc
Matches
                                                                                                                                                                                                                                                                                                                                                                                           RESULT 9
              X888888X8
                                                                                                                                                                                                                                                                                      It is called asS-28 because it is an analogue of anglerfish somatostatin (SS-28). It is more potent than either somatostatin.14 (SS-14) or SS-28 at inhibiting insulin secretion for the treatment of insulinoms. The 14-residue C-terminal peptide is also claimed (aSS-14): aSS-14 is useful for inhibiting insulin secretion by the pancreas aSS-28 and aSS-28 may be useful for decreasing gastric acid secretion and influencing thermoregulation. Their reduced linear forms, wherein the disulphide bridge is not present and is replaced by H, is also claimed. (Updated on 25-MAR-2003 to correct PA field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Growth Hormone secretion; GH; diabetes mellitus; angiopathy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Somatostatin derivs. prodn. - useful for treating diabetes, acromegalia and anglopathy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 83; DB 10; Length 28;
Pred. No. 1e-05;
0; Mismatches 1; Indels
                                                                                                                                                                                        Angler fish somatostatin-28 and fragments - useful in inhibiting insulin secretion and insulinoma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /note= "Cys-pyrrolidine amide"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Location/Qualifiers .
14
                                                             (SALK ) SALK INST BIOLOGICAL STUDIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAP10324 standard; peptide; 14 AA
                                                                                                                                                                                                                                                      Claim 1; page 65; 8pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ő
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 1; Page 6; 8pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 1981-21515D/13 (21515D).
  87US-0033295.
84US-0646610.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Match 93.3%;
Local Similarity 92.9%;
les 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             80CH-0125375.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (updated)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Somatostatin deriv. A28.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  acromegaly; diagnosis.
                                                                                                                                            WPI; 1989-113910/15.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (SANO ) SANDOZ AG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              28 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Key
Modified-site
01-APR-1987;
31-AUG-1984;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 27-FEB-1981.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             11-SEP-1980;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     01-FEB-1984;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     25-MAR-2003
15-DEC-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CH621770-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sandrin E,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local S1
Matches 13;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAP10324;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 8
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õ 셤 Benoit RA;

Ling NC,

Indels

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inserted in a cloning vehicle for expression as such, the gene is preceded by a start codon and immediately followed by one or more termination or stop codons. AAN30112 also exemplifies a further employment, i.e., the provision of cohesive terminal, preferrably comprising one of the two strands of a restriction endonuclease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Polypeptide prodn. by genetically transformed microorganism - in form of cleavable conjugate with another protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (Updated on 16-AUG-2002 to add missing OS field.) (Updated on 25-MAR-2003 to correct PR field.)
     Pred. No. 4.3e-05;
1; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 77; DB 4;
Pred. No. 4.3e-05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The synthetic somatostatin gene is prepd. fragments composed of codons preferred fo genomes. Where the structural gene of a d
                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence of synthetic somatostatin gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic gene; heterologous protein.
                                                                                                                                                                                                                                                 AAP30668 standard; Protein; 14 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Disclosure; Fig 2; 23pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAP40316 standard; Protein; 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 1983-07104K/03 (07104K).
N-PSDB; AAN30312.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 86.5%;
     Similarity 85.7%;
12; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         79US-0090980
                                                                                                                                                                                                                                                                                                                                                                  (updated)
(first entry)
                                                                             1 AGCKNFYWKGFTSC 14
                                                                                                                           1 AGCKNFFWKTFTSC 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 AGCKNFYWKGFTSC 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 AGCKNFFWKTFTSC 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 86.5
Best Local Similarity 85.7
Matchés 12; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                             25-MAR-2003
16-AUG-2002
31-OCT-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         05-NOV-1979;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       26-SEP-1985;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US4366246-A.
                                                                                                                                                                                                                                                                                             AAP30668;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Riggs AD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sednence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAP40316;
        Best Local
Matches 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAP40316
ID AAP4
XX
AC AAP4
                                                                                                                                                                                                                          AAP30668
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         à
                                                                                                                                                                                                                                                                                                                               carbohydrate moleties for incorporation are the pentose and hexose monosaccharides, particularly those having the pyranose structure, such as glucose and fructose.
                                                                                                                                                                                           secretion of glucagon and insilin by the pancreas and secretion of vasoactive intestinal polypeptides, secretin, gastrin. and gastric II has a similar inhibitory efficiency to native somatostatin. Th
                                                                                                                      sequence has an extended biological half-life compared with
                                                                                                                                                                                                                                         It has a similar inhibitory efficiency to native somatostation.
Increased acting inhibitory effect of this peptide is due to the inclusion of a carbohydrate moiety in to the somatostatin chain. This carbohydrate moiety can be linked to either Ser, Asn or Thr which appear in the somatostatin peptide chain. Convenient which appear in the somatostatin peptide chain.
Somatostatin glycosylated analogues - with prolonged half-lives and similar activities and potency to somatostatin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The sequence represents somatostain, which may be expressed after synthetic gene expression in microorganisms. (Updated on 25-MAR-2003 to correct PR field.) (Updated on 25-MAR-2003 to correct PA field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Recombinant microbial cloning vehicle - for expression of polypeptide(s), esp. hormones such as somatostatin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 77; DB 3; Length 14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 77; DB 2; 1
Pred. No. 4.3e-05;
1; Mismatches 1,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAP20165 standard; Protein; 14 AA
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                                                                  Claim 1; Page 11; 13pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 1982-97410E/45 (97410E).
N-PSDB; AAN20160.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity 85.7%;
Matches 12; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 AGCKNFYWKGFTSC 14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   26-SEP-1985;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  05-NOV-1979;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              25-MAR-2003
15-SEP-1992
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AAP20165;

RESULT 10 AAP20165

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Sequence

Matches

Gaps

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; Indels

Sequence Query Match

Length 14;

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ch protein conjugates is
The heterologous protein is
Bacterial cultures comprising
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic gene; somatostatin; cloning vehicle; beta-galactosidase; fusion protein; vaccine; radioimmunoassay;
                                                                                                                                                                                                                                             The inventors claim a transformant bacterial culture cloned from one or more bacteria, each comprising a recombinant microbial vehicle. The vehicle comprises: (requion)-(codons for additional protein) - (codons for desired heterologous polypeptide)-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Recombinant cloning vehicle contg. gene for somatostatin expressed as immunogenic fusion prod. with bacterial polypeptide
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                                                                                                                         Transformant bacterial culture cloned from bacteria - for cultivation to give somatostatin etc. linked to protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 14;
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Pred. No. 4.3e-05;
1; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (Updated on 16-AUG-2002 to add missing OS field.)
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AAP60863
ID AAP60863 standard; protein; 14 AA
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                                                                                                                                                                                                 Disclosure; Fig 2; 23pp; English.
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Best Local Similarity 85.7%;
Matches 12; Conservative
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77US-0849591.
79US-0090980.
85US-0787871.
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(first entry)
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                                                       WPI; 1984-056011/09.
N-PSDB; AAN40292.
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N-PSDB; AAN60778.
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05-JUL-1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       05-NOV-1979;
16-OCT-1985;
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             Riggs AD;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The inventors claim a recombinant microbial cloning vehicle for host expression of polypeptide. Plasmids pSOM 1 and pSOM 11 are new. The Polypeptide is expressed from a synthetic gene, and it is esp. somatostatin e.g. described in US3904594.
                                                                                           Growth hormone inhibitor; insulin inhibitor; glucagon inhibitor; acromegaly therapy; pancreatitis; insulin-dependent diabetes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Recombinant microbial cloning vehicle - esp. useful for host expression of polypeptide e.g. somatostatin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 14;
                                         Sequence encoded by synthetic gene for somatostatin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Protein conjugate; transformed bacteria; plasmid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 86.5%; Score 77; DB 5; 1
Best Local Similarity 85.7%; Pred. No. 4.3e-05;
Matches 12; Conservative 1; Mismatches 1.
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77US-0849591.
79US-0090980.
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77US-0849591,
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(first entry)
11-FEB-1992 (first entry)
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14 AA;

Sequence

30-JUL-1982; 30-JUL-1982; 08-NOV-1977;

14-FEB-1984.

US4431739-A.

Synthetic.

16-AUG-2002 03-AUG-1992

AAP40808;

RESULT 13

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Gaps

WPI; 1984-029658/05. N-PSDB; AAN40257.

Riggs AD;

30-JUL-1982; 08-NOV-1977; 05-NOV-1979;

30-JUL-1982;

10-JAN-1984

Homo sapiens. US4425437-A:

0; - Gaps

Length 14; Indels

Score 77; DB 8; Pred. No. 4.3e-05; 1; Mismatches 1

14 AA;

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Search completed: August 13, 2003, 14:50:59 Job time : 28,2051 secs
                                                Query Match
Best Local Similarity 85.7%;
Matches 12; Conservative
                     Sequence
        X S
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                                                                                                                                                                                                                                ö
       which is present in a
ulon, one or more
                          termination codons, and a DNA sequence encoding an additional protein, especially adjactostidase. Somatostatin is expressed as a fusion protein which can be cleaved which CNBr to give the two proteins. Temporary incorporation of somatostatin into a fusion prod. protects it from in vivo degradation by andogenous enzymes. Alternatively, the fusion protein can be used to raise antibodies for RIA, or in the prepn. of
                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                This sometostatin deriv. has the same activity as natural someto-
statin but has a more specific and longer lasting action. It is
useful in the treatment of diabetes and gastrointestinal disorders
at a dosage of 0.05-0.2 mg/kg.
(Updated on 09-JAN-2003 to add missing OS field.)
(Updated on 25-MAR-2003 to correct PA field.)
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                                                                                                                                                                                                Length 14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New N-acylated somatostatin derivs. - useful for treating diabetes and gastrointestinal disorders
                                                                                                                                                                                            Score 77; DB 7; Length 14;
Pred. No. 4.3e-05;
1; Mismatches 1; Indels
                 regulon,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Somatostatin; diabetes; gastrointestinal disorders:
                                                                                                                        'accines.
(Updated on 31-OCT-2002 to add missing OS field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          label- N-acylated alanine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 |..4
|abel= N-acylated lysine
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/label= N-acylated lysine
3..14
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                                                                                                                                                                                                                                                                                                                                                           AAP70928 standard; protein; 14 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         N-acylated somatostatin deriv.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 1; page 1; 3pp; German.
                                                                                                                                                                                           Query Match
Best Local Similarity 85.7%;
Matches 12; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                          (updated)
(updated)
(first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (DIAL ) DIAMALT AG. (JUNG/) JUNG G.
                                                                                                                                                                   Sequence 14 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Disulfide-bond
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Key
Modified-site
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09-JAN-2003
12-MAR-1991
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Unidentified
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AAP70928
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SUMMARIES

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Description	Segmence 2 Appli	Sequence 2. Appli	2	37	Segmence 52, April	. 2	Segmence 3. Appli		Segmence 53 April	Segmence 8 Appli	Sequence 8. Appli			Segmente 23 April	2
ΩI	US-09-316-505-2	US-10-224-640-2	US-10-251-703-18	US-10-101-487-37	US-10-101-487-52	US-09-766-396-3	US-10-062-375-3	US-09-280-030-64	US-10-101-487-53	US-09-766-396-8	US-10-062-375-8	US-10-221-841-9	US-09-766-396-23	US-10-062-375-23	US-09-766-396-7
jth DB	14 10	14 12	14 12	14 14	15 14	6	14	140 9	14	14 9	14 14	14 15	15 9	15 14	.6 62
Query Match Length DB	86.5	86.5	86.5	86.5	86.5			•		70.8	70.8	. 8 02	70.8	70.8	70.8
Score	77	77	77	77	77	77	77	77	77	.63	63	63	. 63	63	63
Result No.	-	7	m	4	S	9		æ	6	10	11	12	13	14	15

	; APPLICANT:	Elias, Kathleen A. McDowell, Robert S.	· C				
	APPLICANT:	Rawson, Thomas E.					
	APPLICANT	Stanley, Mark S.					
	; TITLE OF II	IITLE OF INVENTION: LOW MOLECULAR WEIGHT PEPTIDOMIMETIC GROWTH HORMONE SECRETAGOG	CULAR WEIGHT	PEPTIDOMIM	TIC GROWT	H HORMONE	SECRETAGOG
	; FILE REFERE	NCE: P0850D2					
	CURRENT AP	CURRENT APPLICATION NUMBER: US/09/316,505	18/09/316,50				
	; CURRENT FIL	CURRENT FILING DATE: 1999-05-21	5-21				
٠.	PRIOR APPL.	PRIOR APPLICATION NUMBER: US 09/057,074 PRIOR FILING DATE: 1998-04-08	09/057,074				
	, NUMBER OF	NUMBER OF SEQ ID NOS: 2			•		
	. ; SEQ ID NO 2	•					
	; LENGTH: 14		٠.				
	; TYPE: PRT						
	ORGANISM: H US-09-316-505-2	ORGANISM: Homosapiens 9-316-505-2					
	Query Match Best Local Similarity	86.58; imilarity 85.78;	Score 77; DB 10; Pred. No. 4.4e-05	DB 10;	Length 14;		
	Matches 12	12; Conservative	7	thes 1;	Indels	0; Gaps	0;
	0y 1	AGCKNFYWKGFTSC 14	,	·.	٠.		
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	RESULT 2						
٠	US-10-224-640-2	2	,				
	; sequence 2, ; Publication	sequence 2, Application US/10224640 Publication No. US20030139348A1	2464U	٠			

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PPLICANT: TOMPKINS, CHRISTOPHER K.
PPLICANT: WAGGONER JR., DAVID W.
TILLE OF INVENTION: RECOMBINANT PRODUCTION OF POLYANIONIC POLYMERS AND USES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GGONER JR., DAVID W.
TION: RECOMBINANT PRODUCTION OF POLYANIONIC POLYMERS AND USES.
TION: THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                , OTHER INFORMATION: Description of Artificial Sequence: Synthetic peptide US-10-101-487-52
                                                                                                                                                                                                                                                                                                                                                                                                CTHER INFORMATION: Description of Artificial Sequence: Recognition US-10-101-487-37
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ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ORGANISM: Artificial Sequence
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Best Local Similarity 85.7
Matches 12; Conservative
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Best Local Similarity 85.7
Matches 12; Conservative
DAVID W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DAVID W.
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                                                                             Stanley, Mark S. VENTION: LOW MOLECULAR WEIGHT PEPTIDOMIMETIC GROWTH HORMONE SECRETAGOGUES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 OTHER INFORMATION: Description of Artificial Sequence: Extracellular of OTHER INFORMATION: Agonist Peptide Sequence US-10-251,703-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 14;
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TILE OF INVENTION: 6 PROCESIN Coupled Receptor Agonists at TILE OF INVENTION: Methods of Activating and Inhibiting ITLE OF INVENTION: Receptors Using the Same
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US/10/251,703
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o. US20030148449A1
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Best Local Similarity 85.7%;
Matches 12; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 AGCKNFFWKTFTSC 14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity 85.7
Matches 12; Conservative
                                                                                                                                                                                                                                                                                                                                                                                           ORGANISM: Homosapiens
'S-10-224-640-2
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APPLICATION NUMBER: US/10/062,375
FILING DATE: 30-Jan-2002
CLASSIFICATION: <UNKNOWN>
                                                                                                                                                                                                               APPLICATION NUMBER: US/08/857,389
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; FRAGMENT TYPE: C-terminal
SEQUENCE DESCRIPTION: SEQ ID NO: 3:
US-10-062-375-3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          64, Application US/09280030A
o. US20010021515A1
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: Kudo, Toshiyuki
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ARLIER APPLICATION NUMBER: JP1
ARLIER FILING DATE: 1998-03-31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity 85.7%;
Matches 12; Conservative
                            TER READABLE FORM:
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12; Conservative
                                                                                                                        CURRENT APPLICATION
                                                                                                                                                                                                PRIOR APPLICATION
                                                                                                                                                                                                                                                                                                                           TELECOMMUNICATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      , NUMBER OF SEQ ID NOS: 6
SOFTWARE: Patentin Ver.
SEQ ID NO 64
                                                                                                                                                                                                                                                      ATTORNEY/AGENT
                                                                                                                                                                                                                                                                                                                                                                            INFORMATION FOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MOLECULE
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Matches
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                                                                                                                                                                                                      EE: THE SCRIPPS RESEARCH INSTITUTE
10666 No. US20020013456Alth Torrey Pines Road, TPC-8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADDRESSEE: THE SCRIPPS RESEARCH INSTITUTE STREET: 10666 No. US20020133000A1th Torrey Pines Road, CITY: La Jolla
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 110;
                                                                                                                                                                                                                                                                                                                                                                                           PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CORTISTATIN: NEUROPEPTIDES
COMPOSITIONS AND METHODS
                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US/09/766,396
FILING DATE: 18-Jan-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1; Mismatches
                                                                                                                                                                                                                                                                                                                                                COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ATION NUMBER: 08/857,389
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:

APPLICANT: Sufcliffe, Gregor J.

de Lecea, Luis

Siggins, George R.

Henriksen, Steven J.

TITLE OF INVENTION: CORTISTATIN
                                                        Sutcliffe, Gregor J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       : <Unknown>
                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
pplication US/09766396
20020013456A1
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Publication No. US20020133000A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            : 110 amino acids
amino acid
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                                                                                                                             TITLE OF INVENTION: CORT
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                                                                                                                                                                                                                                                            California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION DATA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 AGCKNFYWKGFTSC 14
                                                                                                                  riksen,
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Best Local Similarity 85.7
Matches 12: Conservative
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CORRESPONDENCE ADDRES
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                                                                                                                                                                    NUMBER OF
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TILE OF INVENTION: DAMS ENCODING NEW FUSION PROTEINS AND PROCESSES FOR TILE OF INVENTION: PREPARING USEFUL POLYPEPTIDES THROUGH EXPRESSION OF TILE OF INVENTION: DNAS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OTHER INFORMATION: Description of Artificial Sequence: Designated is COTHER INFORMATION: an amino acid sequence of OTHER INFORMATION: MWPSp-WWPmp20-(His)6-EGF-TEV-Somatostatin 28 US-09:280-030-64
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 77; DB 9; Length 140;
Pred. No. 0.00033;
1; Mismatches 1; Indels
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Length 14; ; Indels

us-09-727-739b-2.rapb

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ADDRESSEE: THE SCRIPPS RESEARCH INSTITUTE
STREET: 10666 No. US20020133000Alth Torrey Pines Road, TPC-8
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FRAGMENT TYPE: C-terminal
SEQUENCE DESCRIPTION: SEQ ID NO: 8:
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GENERAL INFORMATION:
APPLICANT: Sutcliffe, Gregor J
de Lecea, Luis
Siggins, George R.
Stggins, George R.
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TITLE OF INVENTION: CORTIST
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STATE: California
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Best Local Similarity 75.0
Matches 9; Conservative
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                                                                                                                                                                            TELECOMMUNICATION
                                                                                                                                                                                                                                                                                                               INFORMATION FOR SEQ ID
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GREGORER A.
GRONER JR., DAVID W.
FTON: RECOMBINAT PRODUCTION OF POLYANIONIC POLYMERS AND USES
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Pred. No. 0.00046;
1; Mismatches 1; Indels
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Sequence 8, Akr.
Patent No. US200200134....
GENERAL INFORMATION:
GENERAL INFORMATIO
                                                                                                                                                                                                                                                           Application US/10101487
o. US20020169125A1
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Siggins, George
Henriksen, Stev
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                                                                                                                                                                                                                                                                                                                                                                                                   G, DAVID W.
GMAN, PHILIP A.
DUIST, ALAN
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Best Local Similarity 85.7%;
Matches 12; Conservative
127 AGCKNFFWKTFTSC 140.
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COMPUTER READABLE FORM:
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US-09-766-396-8
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APPLICANT: FUKUSUMI, Shuji
TITLE OF INVENTION: NO. US20030082648Alel mas-like Receptor Protein and its DNA
TILE REPERBNCE: 2700 USOP
CURRENT APPLICATION NUMBER: US/10/221,841
CURRENT FILIMG DATE: 2002-09-12
PRICOR APPLICATION NUMBER: PCT/JP01/02053
PRICOR FILING DATE: 2001-03-15
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10666 No. US20020013456A1th Torrey Pines Road, TPC-8
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COMPOSITIONS AND METHODS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
; SEQUENCE DESCRIPTION: SEQ ID NO: 8: US-10-062-375-8
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Siggins, George R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               3, Application US/09766396
US20020013456A1
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FILING DATE: 18-Jan-20
CLASSIFICATION: <Unkno
                                                  Query Match
Best Local Similarity 75.0%;
Matches 9; Conservative
                                                                                                                                3 CKNFYWKGFTSC 14
                                                                                                                                                        2 CKNFFWKTFSSC 13
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Best Local Similarity 75.0
Matches 9; Conservative
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STATE: Californ
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GENERAL INFORMATION:
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; ORGANISM: Rat
US-10-221-841-9
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18: THE SCRIPPS RESEARCH INSTITUTE
10666 No. US20020133000Alth Torrey Pines Road, TPC-8
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                                                                                                                                                                                                                                                                                                              Score 63; DB Pred. No. 0.002; Mismatches
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IG DATE: 30-Jan-2002
HEICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US/08/857,389
                                                                                                                                                                                                           MOLECULE TYPE: Protein
FRAGMENT TYPE: C-terminal
SEQUENCE DESCRIPTION: SEQ ID NO: 23:
US-09-766-396-23
NAME: Schmonsees, William REGISTRATION NUMBER: 31,796
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          quence 23, Application US/10062375
blication No. US20020133000A1
GENERAL INFORMATION:
APPLICANT: SUCILIFFE, Gregor
                                                                                                                                                                                                                                                                                                         Query Match 70.8%;
Best Local Similarity 75.0%;
Matches 9; Conservative
                                                                                                                                  SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Siggins, Ger
Henriksen,
                                                                                                                                                                                                                                                                                                                                                                                       3 CKNFYWKGFTSC 14
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                                                      TELECOMMUNICATION
                                                                                                             INFORMATION FOR SEQ
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Gaps
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10666 No. US20020013456Alth Torrey Pines Road,
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; FRAGNENT TYPE: C-terminal
; SEQUENCE DESCRIPTION: SEQ ID NO: 23:
US-10-062-375-23
                                                          Query Match 70.8%; Score 63; Best Local Similarity 75.0%; Pred. No. (Matches 9; Conservative 2; Mismatch
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE DESCRIPTION: SEQ ID NO: 7: US-09-766-396-7
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Best Local Similarity 75.0%;
Matches 9; Conservative
                                                                                                                     3 CKNFYWKGFTSC 14
                                                                                                                                         3 CKNFFWKTFSSC 14
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                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION:
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Search completed: August 13, 2003, 14:53:42 Job time: 13.641 secs

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GenCore version 5.1.6

Copyright (c) 1993 - 2003 Compugen Ltd.

OM protein - protein search, using sw model

Run on: August 13, 2003, 14:47:57; Search time 8.97436 Seconds (without alignments)

150.023 Million cell updates/sec

Title: US-09-727-739B-2
Sequence: 1 AGCKNFYWKGFTSC 14
Sequence: 1 AGCKNFYWKGFTSC 14
Scoring table: BLOSUM62
Gapop 10.0, Gapext 0.5
Searched: 283308 segs, 96168682 residues

Total number of hits satisfying chosen parameters: 283308
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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
Database: PIR.76:*

Minimum DB seq length: 0 Maximum DB seq length: 200000000

Database : PIR_76:*.
1: pir1:*
2: pir2:*
3: pir3:*
4: pir4:*

Pred. No. 1s the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

	Description	somatostatin-25 -	somatostatin II pr	Somatostatin II pr	II	II	-25		somatostatin I - c	somatostatin I - E	somatostatin I - s	somatostatin-28 -	somatostatin-relat	somatostatin I pre	-14	a	somatostatin-14 pr	⊢	somatostatin I pre	somatostatin precu	somatostatin precu	somatostatin precu	somatostatin precu	somatostatin I pre	somatostatin, panc	somatostatin - spo	-14	cortistatin precur	cortistatin-like p	somatostatin-22 pr
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SUMMARIES																														
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dP	Query	100.0	100.0	100.0	100.0	100.0	88.8	86.5	86.5	86.5	86.5	86.5	86.5	86.5	86.5	86.5	86.5	86.5	86.5	86.5	86.5	86.5	86.5	86.5	82.0	80.9	71.9	70.8	67.4	53.9
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("Conlon, J.M.; Davis, M.S.; Falkmer, S.; Thim, L.
T. Blochem. 168, 647-652, 1887
"Title: Structural characterization of peptides derived from prosomatostatins I and Reference number: S00166; MUID:88029486; PMID:2889597

somatostatin II precursor - European flounder (tentative sequence) (fragments) C;Species: Platichthys flesus (European flounder) C;Date: 07-Sep-1990 #sequence_revision 07-Sep-1990 #text_change 31-Mar-2000 C;Accession: S00169

lecule type: protein sidues: 1-10;11-45;46-73 <CON> .e: three peptides which probably originate from a common precursor, were isolat.

C; Keywords: glycoprotein; neuropeptide; pancreatic islet F;1-10/Product: peptide F1 #status experimental <PF1>F;11-45/Product: peptide F3 #status experimental <PF3>F;67-73/Poduct: peptide F2 #status experimental <PF3>F;62-73/Disulfide bonds: #status experimental

Superfamily: somatostatin

30 48 53.9 14 31 47 52.8 42 32 45 50.6 14 34 45 50.6 14 34 45 50.6 14 35 43 48.3 12 36 43 48.3 12 37 43 48.3 12 38 42 47.2 34 39 42 47.2 34 41 46.1 13 41 41 46.1 15 42 41 46.1 15 42 41 46.1 15 43 48.3 15 50matostatin-25 - coho salm 70 Ablerate names: somatost Cipate: 30.Sep-1993 #sequen Cipate: 30.Sep-1993 #s	48 53 45 50 45 50 43 48 43 48 43 48 43 48 41 46 41 46	310 48 53.9 147 2 32 45 50.6 147 2 34 45 50.6 147 2 34 45 50.6 147 2 34 45 50.6 147 2 35 43 48.3 139 2 36 43 48.3 139 2 37 43 48.3 139 2 40 41 46.1 139 2 41 46.1 139 2 41 46.1 139 2 42 47.2 346 2 43 48.3 199 2 44 41 46.1 139 2 44 41 46.1 580 2 45 41 46.1 580 2 46 41 46.1 580 2 47 41 46.1 580 2 48 41 46.1 580 2 49 41 46.1 580 2 40 61 580 580 2 41 46.1 580 580 2 42 47.2 580 2 43 48.3 772 3 44 41 46.1 580 2 45 41 56.1 580 2 45 41 66.1 580 2 40 62 62 62 62 62 62 62 62 62 62 62 62 62	42.01.20 (1.14.4.3.1.1.3.1.1.3.1.1.3.1.1.3.1.1.3.1.1.3.1.1.3.1		00864 66275 66275 66275 1087 11847 11847 11847 1184 1184 1189 1189 1189 1189 1189 1189	MEN SSOI SOI SOI SOI SOI SOI SOI SOI SOI SO	m // / / / / / / / / / / / / / / / / /	tt_che kisuu) kisuu) kisuu) kisuu) kisuu odani> dani> nogth indel		hetical prote hetical prote hetical prote hetical prote hetical prote oxide dismuta ide channel p sterone-resis hasmic reticulate nastric reticulate nastric reticulate oxyrobiaes oxyrosidase [1 hypothetical hypothetical islet somatos; ut rather ded	. And Eatin	is is
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Molecule type: mRNA

Stasiduse: 1.125 - 4003-

(Cross references: GB:V00641; GB:J00947; GB:M23199; NID:964030; PIDN:CAA23987.1; PII

Experimental source: 1slet tissue (endocrine pancreas)

Spiess, J.; Noe, B.D.

171 - 171 - Acad. Sci. U.S.A. 82, 277-281, 1985

Title: Processing of an anglerfish somatostatin precursor to a hydroxylysine-conta.

171 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 172 - 17
                                                                                                                                              Date: 31-Mar-1981 #sequence_revision 31-Mar-1981 #text_change 28-May-1999
Accession: B93236, A94038; A27376; A01434; A21881; A93236
Abobart, P.; Crawford, R.; Shen, L.; Pictet, R.; Rutter, W.J.
Atture 288, 137-141, 1980
Title: Cloning and sequence analysis of cDNAs encoding two distinct somatostatin preference number: A93236; MUID:81052423; PMID:6107860
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A;Molecule type: protein
A;Residues: 98-125 <RPI>
A;Andreway. P.C.; Nichols. R.; Dixon, J.E.
J. Biol. Chem. 262, 12692-12699, 1987
A;Title: Post-translational processing of preprosomatostatin-II examined using fast
A;Reference number: A27376; MUID:87308304; PMID:2887572
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A; Residues: 17-76, DV, 79-89, 'G', 91-125 <AND>
A; Residues: 17-76, DV, 79-89, 'G', 91-125 <AND>
A; Residues: 17-76, DV, 79-89, 'G', 91-125 <AND>
C; Superfamily: somatostatin
C; Superfamily: somatostatin; neuropeptide; pyroglutamic acid
E; 12-24/Domain: stopeptide $status experimental <SIG>
F; 25-97/Domain: propeptide $status experimental <RRO>
F; 97-125/Product: somatostatin II $status experimental <AND>
F; 17-125/Product: somatostatin II $status experimental <AND>
F; 114-125/Disulfide bonds: $status experimental
F; 120/Modified site: hydroxlyshe (Lys) $status experimental
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R;Collon, J.M.; Deacon, C.F.; Hazon, N.; Henderson, I.W.; Thim, L.

Gen. Comp. Endocrinol. 72, 181-189, 1989

A;Title: Somatostatin-related and glucagon-related peptides with unusual structural

A;Reference number: A60840; MUID:89065329; PMID:2904391
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              C;Species: Anguilla anguilla (European eel)
C;Date: 30-Sep-1993 #sequence_revision 30-Sep-1993 #text_change 20-Mar-1998
C;Accession: B60840
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A; Molecule type: protein
A; Residues: 1.25 <CON>
C; Superfamily: somatostatin
C; Keywords: hormone; hydroxylysine; pancreatic islet
F; 20/Modified site: 5-hydroxylysine (Lys) (partial) #status experimental
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 89; DB 1; Length 125; Pred. No. 6.2e-07;
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Pred. No. 5e-06;
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92.9%;
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;Date: 13-Sep-1996 #sequence_revialnbow trout)

;Accession: 151064

R;Moore, C.A.; Kittilson, J.D.; Dahl, S.K.; Sheridan, M.A.

Gen. Comp. Endocrinol. 98, 253-261, 1995

A;Title: Isolation and characterization of a cDNA encoding for preprosomatostatin contain A; Reference number: 151064; MUID:95354921; PMID:7628684
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          somatostatin II precursor - shorthorn sculpin (tentative sequence) (fragments)
C;Species: Wyoxocephalus scorpius (shorthorn sculpin, daddy sculpin)
C;Date: 07-Sep-1990 #sequence_revision 26-Jan-1996 #text_change 31-Mar-2000
C;Accession: S00166; A26993
Eur. J. Biochion, J.M.; Davis, M.S.; Falkmer, S.; Thim, L.
Eur. J. Biochem. 168, 647-652, 1987
A;Ttile: Structural characterization of peptides derived from prosomatostatins I and II
A;Reference number: S00166; MUID:88029486; PMID:2889597
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   common precursor, were isolated
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                                                                                                Gaps
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Best Local Similarity 100.0%; Pred. No. 3.8e-07;
Matches 14; Conservative 0; Mismatches 0;
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          Score 89; DB 2;
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Ouery Match 100.0%;
Best Local Similarity 100.0%;
Matches 14; Conservative 0;
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R;Spiess, J.; Villarreal, J.; Vale, W.

Biochemistry 20, 1982-1988; 1981

A;Title: Isolation and sequence analysis of a somatostatin-like polypeptide from ovi)

A;Reference number: A61322; WUID:81184502; PMID:7225368

A;Accession: A61322
                                                                                                                                                                                                                                                                                                           R'Burgus, R.; Ling, N.; Butcher, M.; Guillemin, R.
Proc. Neull, Acad. Sci. U.S.A. 70, 684-688, 1973
A;Title: Primary structure of somatostatin, a hypothalamic peptide that inhibits the
A;Reference number: A61344; MUID:7320956; PMID:4314982
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             somatostatin-related protein - Atlantic hagfish
C;Species: Myxine glutinosa (Atlantic hagfish)
C;Date: 21-May-1990 #sequence_revision 21-May-1990 #text_change 31-Dec-1993
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C;Reywords: neuropeptide
C;Reywords: somatostatin-28 #status experimental <528>
F;11-28/Product: somatostatin-14 #status experimental <514>
F;17-28/Disulfide bonds: #status experimental
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Pred. No. 1.1e-05;
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6e-06;
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A Residues: 1-14 <CON>
A Note: the source is designated as Cottus scorpius
C; Superfamily: somatostatin
C; Keywords: neuropeptide; pancreatic islet
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1; Mismatches
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llarity 85.7%; Pr
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nes 12; Conservative
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A; Residues: 1-28 <SPI>
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Peptides 11, 461-466, 1990
Peptides 11, 461-466, 1990
A:Title: Isolation and structural characterization of insulin, glucagon and somatostatin A:Reference number: A60414; MUD:90341082; PMID:1974347
A:Retession: C60414
A:Status: preliminary
A:Molecule type: protein
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A:Molecule type: protein
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A;Title: Somatostatin-related and glucagon-related peptides with unusual structural feat A;Reference number: A60840; MUID:89065329; PMID:2904391
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Cippedies: Oncortyprochus Kisutch (coho salmon)
Cipate: 30-Sep-1993 #sequence_revision 30-Sep-1993 #text_change 07-May-1999
Cipate: 30-Sep-1993 #sequence_revision 30-Sep-1993 #text_change 07-May-1999
Riplisetskaya, E.M.; Pollock, H.G.; Rouse, J.B.; Hamilton, J.W.; Kimmel, J.R.; Andfews,
Gen. Comp. Endocrinol. 63, 252-263, 1986
A;Tille: Characterization of Coho salmon (Oncorhynchus kisutch) islet somatostatins.
A;Reference number: A60842; MUID:87055212; PMID:2877919
                                                                                                                          Species: Pseudemys scripta (slider)
Date: 11-Feb-1993 #sequence_revision 11-Feb-1993 #text_change 31-Dec-1993
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Date: 30-Sep.1993 #sequence_revision 30-Sep.1993 #text_change 20-Mar-1998
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C; Superfamily: Somatostatin
C; Keywords: hormone; pancreatic islet
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A; Accession: A60840
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RESULT 7
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Nicteraturus (channel caffish)
Nicresian: S00292; MUID: 82265698; PMID: 6179939
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Ryaylor, W.L.; Oollter, K.J.; Deschenes, R.J.; Welth, H.L.; Dixon, J.E.
Proc. Natl. Acad. Sci. U.S.A. 78, 6694-6698, 1981
A;Rtile: Sequence analysis of a cDNA coding for a pancreatic precursor to somatostat A;Reference number: A93897; MUID:82082515; PMID:6171821
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A; Residues: 82-108 < TAXY
A; Cross-references: GB:J00944
B; Andrews, P.C.; Dixon, J.E.
J. Biol. Chem. 256, 8267-8270, 1981
A; Title: Isolation and structure of a peptide hormone predicted from a mRNA sequence
A; Reference number: A92334; MUID:81264223; PMID:6114953
A; Accession: A92334
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(S.Species: Id-allurus punctatus (channel catfish)
(S.Species: IJ-Sep-1996 #sequence_revision I3-Sep-1996 #text_change 16-Jul-1999
(S.Accession: ISO798
R:Dixon, J.E.; Andrews, P.C.
Adv. Exp. Med. Biol. 1889, 19-29, 1985
C;Comment: Somatostatin inhibits the release of somatotropin. C;Superfamily: somatostatin (S;Neywords: hormone; hypothalamus; intestine; neuropeptide F;1-64/Domain: propeptide #status experimental -RRO> F;5-92/Product: somatostatin-28 #status experimental -KMS> F;37-92/Product: somatostatin-14 #status experimental -KM4> F;37/Binding site: carbohydrate (Asn) (covalent) #status absent F;81-22/Pusulifide bonds: #status experimental
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A; Residues: 101-114 < AND>
C; Superfamily: somatostatin
C; Superfamily: somatostatin
C; Reywords: neuropeptide
F; 1-24/Domain: signal sequence #status predicted < SIG>
F; 25-100/Domain: propeptide #status predicted < PRO>
F; 101-114/Product: somatostatin: 14 #status experimental < WAT>
F; 103-114/Disulfide bonds: #status experimental
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85.7%; Pred. No. 3.5e-05;
tive 1; Mismatches 1;
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                 R;Conlon, J.M.; Askensten, U.; Falkmer, S.; Thim, L.
Endocrinology 122, 1855-1859, 1988
A;Title: Primary structures of somatostatins from the islet organ of the hagfish suggest
A;Reference number: A32271; MUID:88195948; PMID:2896118
A;Accession: A32271; MUID:88195948; PMID:2896118
A;Recession: A32271
A;Realdues: 1-3 4 <CON>
C;Superfamily: somatostatin
C;Keywords: neuropeptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Nalternate names: prosomatostatin
Nichternate names: prosomatostatin 14 (SS-14); somatostatin 28 (SS-28)
C;Date: 30-Nov-1980 #sequence_revision 31-Jan-1997 #text_change 31-Jan-1997
C;Accession: A34109; A34212; A91273; A91844; A90389; S13616; A01432
Nichternate name: A4109; Name: P.G.A.; Holst, J.J.
Nichter prosomatostatin 1-64 is a major product of somatostatin gene expression in pance Nichternate names: A34109; MUD: 89278131; PMID: 2567292
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A; Residues: 65-92 <8CR.
A; Experimental source: hypothalamus
A; Schally, A.V.; Dupont, A.; Arimura, A.; Redding, T.W.; Nishi, N.; Linthicum, G.L.; Sch
Biochemistry 15, 509-514, 1976
A; Title: Isolation and structure of somatostatin from porcine hypothalami.
A; Reference number: A90398; MUID:76136331; PMID:1252409
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A;Molecule type: protein
A;Residues: 1-64 GERS-
K;Schmidt, W.E.; Mutt. V; Kratzin, H.; Carlquist, M.; Conlon, J.M.; Creutzfeldt, W.
FEBS Lett. 192, 141-146, 1985
A;Title: Isolation and characterization of proSS1-32, a peptide derived from the N-termi
A;Reference number: A24222; MUID:86030691; PMID:2865169
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A;Residues: 65-92 <PRA>
A;Residues: 65-92 <PRA>
A;Residues: 65-92 <PRA>
A;Experimental source: intestine
B;Schally, A.V.; Buang, W.Y.; Chang, R.C.C.; Arimura, A.; Redding, T.W.; Millar, R.P.; F
Proc. Natl. Acad. Sci. U.S.A. 77, 4489-4493; 1980
A;Title: Isolation and structure of pro-somatostatin: a putative somatostatin precursor
A;Reference number: A93854; MUID:81054799; PMID:6107906
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A; Molecule type: protein
A; Residues: 1-32 <503>
B; Residues: 1-10; 55-58, 1980
A; Tille: N-terminally extended somatostatin: the primary structure of somatostatin-28.
A; Reference number: A91273; MUID:80113258; PMID:7353633
A; Reference number: A91273
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A; Residues: 79-92 <SC2>
A; Experimental source: hypothalamus
R; Bersani, M; Johnsen, A.H.; Holst, J.J.
PEBS Lett. 279, 237-239, 1991
A; Title: Oxidation/reduction explains heterogeneity of pancreatic somatostatin.
A; Reference number: S13616; MUID:91160722; PMID:1672110
                                                                                                                                                                                                                                                                                                                                                                                                            Length 34;
                                                                                                                                                                                                                                                                                                                                                                                               Watch 86.5%; Score 77; DB 2; 1 Local Similarity 85.7%; Pred. No. 1.4e-05; les 12; Conservative 1; Mismatches 1.
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Somatostatins of the channel catfish.

A;Status: preliminary; translated from GB/EMBL/DDBJ
A;Molecule type: mRNA
A;Residues: 1-114 < CIN:
A;Cross-references: GB:MZ5903; NID:g213339; PIDN:AAA49339.1; PID:g213340
C;Superfamily: somatostatin Score 77; DB 2; Length 114; Pred. No. 4.3e-05; 1; Mismatches 1; Indels Ouery Match 86.5%; Best Local Similarity 85.7%; Matches 12; Conservative

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0; Gaps

Search completed: August 13, 2003, 14:51:51 Job time : 9.97436 secs

GenCore version 5.1.6 Copyright (c) 1993 - 2003 Compugen Ltd.

OM protein - protein search, using sw model

August 13, 2003, 14:47:17; Search time 5.38462 Seconds (Without alignments) 122.269 Million cell updates/sec Run on:

1 AGCKNFYWKGFISC 14 US-09-727-739B-2 89 Title: Perfect score: Sequence:

Scoring table: BLOSUM62 Gapop 10.0 , Gapext 0.5

127863 seqs, 47026705 residues Searched:

127863 Total number of hits satisfying chosen parameters:

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

SwissProt_41:* Database :

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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100.0%; Score 89; DB 1; Length 73; 100.0%; Pred. No. 3.1e-07; Live 0; Mismatches 0; Indels

Ouery Match Best Local Similarity 100.0 Matches 14; Conservative

7989 MW;

PEPTIDE PEPTIDE DISULFID SEQUENCE

[TYR21,GLY24]SOMATOSTATIN-28. [TYR7,GLY10]SOMATOSTATIN-14. CCCBA6B30DCB29BB CRC64; 01-MAR-1989 (Rel. 10, Created)
01-FBB-1991 (Rel. 17, Last sequence update)
16-072-2001 (Rel. 40, Last annotation update)
Somatostatin II precursor (Contains: [Tyr21,Gly24]somatostatin-28;
[Tyr7,Gly10]somatostatin-14] (Fragments).
Myoxocephalus scorpius (Shorthorn sculpin) (Daddy sculpin).

74 AA.

PRT;

STANDARD;

SMS2_MYOSC

SMS2_MYOSC

RESULT 2

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]son a; E ei; rone	ed from increatic and the EE OF SOP MILY.
	•	1y24 brat eost Pleu	ived from pancreatic in and the fast of SOMP FAMILY.
		te) date) ral(dy24]sc vertebrata; Vertebrata; pha; Pleuron thys.	Hancreas; 880.29486; PubMed=2889597; M., Davis M.S., Fallmer S., Thim L.; rai characterization of peptides derived from costatins I and II isolated from the pancreati of teleostean fish: the daddy sculpin and the Blochem. 168:647-652(1987). TION: SOMATOSTATIN INHIBITS THE RELEASE OF SO ELLUAR LOCATION: Secreted. LARITY: BELONGS TO THE SOMATOSTATIN FAMILY. 169; S00169. on pair of basic residues; Hormone; Multigen 1
Ожықкый нечт	ALIGNMENTS	cr; 73 AA. nnce update) nnce update) ins: [Tyr2].("agments) "anides, Verte Perconorpha, Platichthys	rs., Thim L.; peptides derited from the perides derited from the p. daddy sculpin BITS THE RELEATED. SOMATOSTATIN F.
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	, ž.,.	STANDARD; PR 11. 18, Created) 11. 40, Last seque 11. 40, Last annot 12 precursor [Conta natostatin-14] [Fr sus, European flo rzoa, Chordata Cr Neoptery911; Tel Acanthoptery911; ; Pleuronectidae;);	ESQUENCE. ISSUE-Pancreas; EDILNE-88029486; PubMed-2889597; EDILNE-88029486; PubMed-2889597; Onlon J. M., Davis M.S., Falkmer Structural characterization of prosomatostatins I and II isolate pecies of teleostean fish: the dur. J. Blochem. 168:647-652(1987) I- FUNCTION: SOMATOSTATIN INHIBIT STMICARITY: BELONGS TO THE SO IR; SO0169; SO0169. IR; SO0169; SO0169. I leavage on pair of basic residue Nu TER.
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	ģ	SMS2_PLAFE 927780; 01-MAY-1991 01-MAY-1991 01-MAY-1991 16-CCF-2001 Somatostati [Tyr7, G1y10 [Tyr7, G1y10 Platithhys Eukaryota; Eukaryota; Acathomorp Pletromecto	SEQUENCE. TISSUE-Pa MEDLINE-B CONION J. "Structur prosomato species o Eur. J. B Eur. J
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Moore C.A., Kittison J.D., Dahl S.K., Sheridan M.A.;

Moore C.A., Kittison J.D., Dahl S.K., Sheridan M.A.;

Isolation and characterization of a cDNA encoding for
preprosomatostatin containing (Tyr7, Gly10)-somatostatin-14 from the
endocrine pancreas of rainbow trout, Oncorhynchus mykiss.";
endocrine pancreas of rainbow trout, Oncorhynchus mykiss.";
e. Concornation. 98:23-261(1995).
el- FUNCTION: SOMATOSTAIN INHIBITS THE RELEASE OF SOMATOTROPIN.
el- SUBCELLULAR LOCATION: Secreted.
                                                                                                                           Conlon J.M., Davis M.S., Falkmer S., Thim L.;
"Structural characterization of peptides derived from
yeosomatostatins I and II isolated from the pancreatic islets of two
species of teleostean fish: the daddy sculpin and the flounder.";
Eur. J. Blochem. 168:647-652(1987).
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16-0CT-2001 (Rel. 40, Last sequence update)
16-0CT-2001 (Rel. 40, Last annotation update)
Somatostatin II precursor [Contains: [Tyr21,Gly24]somatostatin-28;
   Vertebrata; Euteleostomi;
Euteleostei; Neoteleostei
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          [Tyr7,Gly10] somatostatin-14].
Oncorhynchus mykiss (Rainbow trout) (Salmo gairdneri).
Bukaryota: Marazoa: Chordata: Craniata: Vertebrata: Euteleostomi; Actinopterygi; Meopterygi; Teleostel; Euteleostel; Protacanthopterygii; Salmoniformes; Salmonidae; Oncorhynchus.
                                                                                                                                                                                                                                                                           MEDLINE-87190954; PubMed-2883025; Cuffield S.M., Carne A., Cuffield J.F.; The amino-acid sequences of sculpin islet somatostatin-28 and peptide YY.";
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               Actinopterygil; Neopterygil; Teleostei; Euteleostei; Neoteleo
Acanthomorpha; Acanthopterygil; Percomorpha; Scorpaeniformes;
Cottoidel; Octildae; Myoxocephalus.
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                                                                                                                                                                                                                                                                                                                                                                                                                             [TYR21, GLY24]SOMATOSTATIN-28.
[TYR7, GLY10]SOMATOSTATIN-14.
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the European Bioinformatics Institute.
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74 AA;
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This SWISS-PROT entry is copyright. It is produced through a collaboration between the SWAiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way
tong as its content is in no way noved. Usage by and for commercial (See http://www.isb-sib.ch/announce/
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                                                                                                                                                                                                                       Pfan; PF03002; Somatostatin; 1.
Cleavage on pair of basic residues; Hormone; Signal; Multigene family.
                                                                                                                                                                                                                                                                                                                                              TYR21_GLY24]SOMATOSTATIN-28 (POTENTIAL).
TYR7, GLY10]SOMATOSTATIN-14.
$Y SIMILARITY
$2.0595022FCA6D91 CRC64;
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MEDLINE—87308304; PubMed=2887572;
Andrews P. C., Nichols R., Dixon J.E.;
Post-translational processing of preprosomatostatin-II examined using fast atom bombardment mass spectrometry.";
Dislol. Chem. 262:12692-12699(1987).
-I- FUNCTION: SOMATOSTAIN INHIBITS THE RELEASE OF SONATOTROPIN.
-I- SUBCELLAREOGS: SOMATOSTAIN INHIBITS.
-I- MISCELLAREOGS: SOMATOSTAIN II MAX HAVE A DIFFERENT DEGREE OF ACTIVITY OR A DIFFERENT TYPE OF TARGET CELL FROM SOWATOSTATIN I.
-I- SIMILARITY: BELONGS TO THE SOMATOSTATIN FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Lopnius americanus (American goosefish) (Anglerfish).
Eukaryota, Metazoa, Chordata, Craniata, Vertebrata; Euteleostomi;
Actinopterygii, Neopterygii, Teleostei, Euteleostei, Neoteleostei,
Acarthomorpha; Paracanthopterygii; Lophiiformes; Lophiidae; Lophius,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Cleavage on pair of basic residues; Hormone; Signal; Hydroxylation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                [Contains: [Tyr7,Gly10]somatostatin-14].
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE FROM N.A. MEDLINE-81053423; PubMed-6107860; MEDLINE-81053423; PubMed-6107860; Flotet R., Rutter W.J.; Hobart P.M., Crawford R., Shen L., Pictet R., Rutter W.J.; "Cloning and sequence analysis of conas encoding two distinct somatostatin precursors found in the endocrine pancreas of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 115;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 89; DB 1; Length 1. Pred. No. 4.8e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Usage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SMS2_LOPAM STANDARD; PRT; 125 AA. P01170; 091066; 21-UUL-1986 (Rel. 01, Created) 01-CCT-1989 (Rel. 12, Last sequence update) 16-CCT-2001 (Rel. 40, Last annotation update) somatostatin II precursor [Contains: [Ty77,61]
                                  removed.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            use by non-profit institutions as long a
modified and this statement is not removed.
entities requires a license agreement (See l
or send an email to license@isb.sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          100.0%; Score 89; Di
11arity 100.0%; Pred. No. 4.6
Conservative 0; Mismatches
use by non-profit institutions as lo
modified and this statement is not remo-
entities requires a license agreement (
or send an email to license@isb-sib.ch)
                                                                                                                                                                                                                                                                                                                                 POTENTIAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PARTIAL SEQUENCE, AND HYDROXYLATION.
                                                                                                                                                         PIR; 151064, 151064
InterPro; IPR004250; Somatostatin.
Pfam; PF03002; Somatostatin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           InterPro; IPR004250; Somatostatin.
                                                                                                                                                                                                                                                                                                                                                     115 [1
115 [1
115 B
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    102 AGCKNFYWKGFTSC 115
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 AGCKNFYWKGFTSC 14
                                                                                                                                            EMBL; U32471; AAC59695.1;
PIR; I51064; I51064.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Nature 288:137-141(1980).
                                                                                                                                                                                                                                                                                                                                                     88
102
104
115 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity
es 14; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                               DISULFID
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Best Local S
Matches 14
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DISULFID

SHEEREES

CONFLICT CONFLICT

SEQUENCE Query Match

Best Loca Matches

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Oreochromis niloticus (Nile tilapia) (Tilapia nilotica).
Mistaryota, Metacoa, Chordata; Cranlata; Vertebrata; Euteleostomi;
Actinopterygii, Neopterygii; Teleostei; Buteleostei; Neoteleostei; Neopterygii; Teleostei; Percomorpha; Acanthopterygii; Percomorpha; Acanthopterygii; Percomorpha; Perciformes; Labroidei;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Nguyen T.M., Wright J.R. Jr., Nielsen P.F., Conlon J.M.; Characterization of the pancreatic hormones from the Brockmann body of the tilapia: implications for islet xenograft studies."; Comp. Biochen. Physiol. 11C:33-44(1995).

- FUNCTION: SOMATOSPATIN INHIBITS THE RELEASE OF SOMATOTROPIN.
-I. SUBCELLULAR LOCATION: Secreted.
-I. SIMILARITY: BELONGS TO THE SOMATOSTATIN FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conlon J.M., Davis M.S., Falkmer S., Thim L.;

Structural characterization of peptides derived from prosomatostatins I and II isolated from the pancreatic islets of two
                                                                                                                                                                                                                         03-NOV-1997 (Rel. 35, Created)
01-NOV-1997 (Rel. 35, Last sequence update)
11-OCT-2001 (Rel. 40, Last amontation update)
16-OCT-2001 (Rel. 40, Last amontation update)
16-OCT-2001 (To recursor [Contains: [Tyr21,Gly24]somatostatin-28;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Myoxocephalus scorplus (Shorthorn sculpin) (Daddy sculpin), Oncorhynchus kisutch (Coho salmon), and Anguilla anguilla (European freshwater eel).

Eukaryota; Metazoa; Chordata; Cranlata; Vertebrata; Euteleostomi; Acthopterygli, Neopterygli, Teleostei, Euteleostei; Rocteleostei, Acanthopterygli, Percomorpha; Scorpaeniformes;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                [TYR21, GLY24]SOMATOSTATIN-28. [TYR7, GLY10]SOMATOSTATIN-14.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 28;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             47C049F4866EF4AC CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 83; DB 1; 1
Pred. No. 1.1e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      01-FEB-1991 (Rel. 17, Created)
01-FEB-1991 (Rel. 17, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
                                                                                                                                                                                                       28 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
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                                                                                                                                                                                                       PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Cottoidei; Cottidae; Myoxocephalus:
NCBI_TaxID=8097, 8019, 7936;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MEDLINE-95384941; PubMed-7656183;
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                                                                       108 GCKNFYWKGFTSC 120
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            15 AGCKNFYWKGLTSC 28
                                             2 GCKNFYWKGFTSC 14
13; Conservative
                                                                                                                                                                                                       STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           Oreochromis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity
es 13; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NCBI_TaxID=8128;
                                                                                                                                                                                                       ORENI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Somatostatin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SMS1_MYOSC
P20750;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DISULFID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
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Matches
Matches
                                                                                                                                                            RESULT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the Exab. Outstation the Exab. Outstation. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed usage by and for commercial entities requires a license agreement (see http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           basic residues; Hormone; Signal; Multigene family.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Lin X.-W., Peter R.E.;
Cloning and characterization of cDNAs encoding preprosomatostatin-I
and -II from goldfish brain.";
Submitted (JUN-1996) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                [Tyr7,Gly10]somatostatin-14].
Gazassius auratus (Goldfish).
Eukaryota: Metazoa: Chordata; Craniata; Vertebrata; Euteleostomi;
Actinopterygil; Neopterygil; Teleostel; Ostarlophysi; Cypriniformes;
Cyprinidae; Carassius.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Jyyuaw, vyrius,
16-OCT-2001 (Rel. 40, Created)
16-OCT-2001 (Rel. 40, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
Somatostatin II precursor [Contains: [Tyr21,Gly24]somatostatin-28;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Otto C.J., Peter R.E.;
The expression of SEP.;
Submitted (SEP-1997) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                       ö
                                                                                                                                                                                                                         Length 125;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYR21,GLY24]SOMATOSTATIN-28.
TYR7,GLY10]SOMATOSTATIN-14.
SY SIMILARITY.
                                                          [TYR7,GLY10]SOMATOSTATIN-14
                                                                                                  HYDROXYLATION.

DV -> TG (IN REF. 1).

G -> E (IN REF. 1).

5E14605D7B9A46FE CRC64;
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98957D68011A651A CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FUNCTION: SOURCES SUBCELULAR LOCATION: Secreted, SUBCELLULAR LOCATION: SECRETED SUBJECT THE SOURCESTATIN FAMILY.
                                                                                                                                                                                                                         Score 89; DB 1; I
Pred. No. 5.2e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 1;
2e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           120 AA
                                                                                                                                                                                                                                                                       Mismatches
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Pred. No.
               POTENTIAL.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Interpro; IPR004250; Somatostatin.
                                                                                                                                                                                                                                                                       ő
               24 P
109
125 [7]
120 H
78 D
90 G
14052 MW;
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100.0%;
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                                                                                                                                                                                                                 Match 100.0%;
Local Similarity 100.0%;
Les 14; Conservative 0
                                                                                                                                                                                                                                                                                                                                      112 AGCKNFYWKGFTSC 125
                                                                                                                                                                                                                                                                                                            1 AGCKNFYWKGFISC 14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           leavage on pair of
                                                                                                                                                                       125 AA;
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NCBI_TaxID-7957;
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ISSUE-Brain

SMS2_CARAU

RESULT 5

ISSUE-Liver

CONFLICT

SULFID

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InterPro; IPR004250; Somatostatin. Pfam; PF03002; Somatostatin; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDLINE-88195948; PubMed-2896118;
                                                                                                                                                                                                                                      1 AGCKNFYWKGFTSC 14
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                                                                                                                                                                                          Conservative
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                                                                     3
14 AA;
                                                                                                                                                              Local Similarity
es 12; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Best Local Similarity
Matches 12; Conser
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SMS_MYXGL
                                                                           DISULFID
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                                                                                                                                           Query Match
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                                                                                                                                                                                                                                                                                                                                                                                    SMS1_AMICA
                                                                                                                                                                    Best Loca
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           "Isolation and structural characterization of insulin, glucagon and somatostatin from the turtle, Pseudemys scripta.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Alligator mississippiensis (American alligator), and
Trachemys scripta (Red-eared slider turtle) (Pseudemys scripta).
Eukaryota; Metazoa; Chordata; Craniata, Vertebrata; Euteleostomi;
Archosauria; Crocodylidae; Alligatorinae; Alligator.
NCBI_TaxID=8496, 34903;
species of teleostean fish: the daddy sculpin and the flounder.";
Eur. J. Biochem. 168:647-652(1987).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FUNCTION: SOMATOSTATIN INHIBITS THE RELEASE OF SOMATOTROPIN. SUBCELLULAR LOCATION: Secreted. SIMILARIT: BELONGS TO THE SOMATOSTATIN FAMILY. 5. C60414; C60414.
                                                                                                                                                                                                                                                                                                                                                        conlon J.M., Deacon C.F., Hazon N., Henderson I.W., Thim L.; Somatostatin-related and glucagon-related peptides with unusual Structural features from the Buropean eel (Anguilla anguilla).*;
                                                                                                                                                                                                                                                                                                                                                                                                                           Gen. Comp. Endocrinol. 72:181-189(1988).
                                                                                      SPECIES-O.ktsutch, TISSUE-Pancreas;
MEDLINE-67055212; PubMed-2877919;
Plisetskay E.M., Pollock H.G., Rouse J.B., Hamilton J.W.,
Kimmel J.N., Andrews P.C., Gorbman A.;
"Characterization of coho salmon (Oncorhynchus kisutch) islet
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 14;
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SIMILARITY: BELONGS TO THE SOMATOSTATIN FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Match 86.5%; Score 77; DB 1; I Local Similarity 85.7%; Pred. No. 4.5e-06; les 12; Conservative 1; Mismatches 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            -UTL-1993 (Rel. 26, Created)
-UTL-1993 (Rel. 26, Last sequence update)
-CCT-2001 (Rel. 40, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  14 AA.
                                                                                                                                                                                                                somatostatins.";
Gen. Comp. Endocrinol. 63:252-263(1986).
                                                                                                                                                                                                                                                                                                            SPECIES-A.anguilla; TISSUE-Pancreas;
MEDLINE-89065329; PubMed-2904391;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          InterPro; IPR004250; Somatostatin.
Pfam; PF03002; Somatostatin; 1.
Hormone; Multigene family.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SPECIES-T.scripta;
MEDLINE-90341082; PubMed-1974347;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        14 AA; 1640 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 AGCKNFFWKTFTSC 14
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Prosomatostatin.I is processed to somatostatin-26 and somatostatin-14 in the pancreas of the bowfin, Amia calva.";
legul. Pept. 47:33-39(1923).
                                                                                                                                                                                                                                                                                                                                                         16-007-2001 (Rel. 40, Created)
16-007-2001 (Rel. 40, Last sequence update)
16-007-2001 (Rel. 40, Last annotation update)
Somatostatin I precursor [Contains: Somatostatin 26; Somatostatin-14]
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       01-NOV-1990 (Rel. 16, Created)
01-NOV-1990 (Rel. 16, Last sequence update)
16-CCT-2001 (Rel. 40, Last annotation update)
Somatostatin precursor [Contains: Somatostatin-34; Somatostatin-14]
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Eukaryota, Metazoa, Chordata, Craniata, Hyperotreti, Myxiniformes,
Myxinidae, Myxininae, Myxine.
                                                                                                                                                                                                                                                                                                                                                                                                                                                        (Fragment).
Amia calva (Bowfin).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Actinopterygii; Neopterygii; Amiiformes; Amiidae; Amia.
NCBI_TaxID=7924;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            -1- FUNCTION: SOMATOSTATIN INHIBITS THE RELEASE OF SOMATOTROPIN.
-1- SINDCELLULAR LOCATION: Secreted.
-1- SIMILARITY: BELLOS TO THE SOMATOSTATIN FAMILY.
-1- IRRO04250; Somatostatin.
-1- FO3002; Somatostatin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         basic residues; Hormone; Multigene family
                                                             Length 14;
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BY SIMILARITY.
D6270F5C09682679 CRC64;
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                                                          Score 77; DB 1;
Pred. No. 4.5e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 77; DB 1;
Pred. No. 8.1e-06;
1; Mismatches 1
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SOMATOSTATIN-14.
                                                                                                                                                                                                                                                                                                                   26 AA.
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                                                                                       ed. No. 4.56
Mismatches
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MEDLINE=94023232; PubMed=8105513;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   26 AA; 2931 MW;
                                                               85.78;
  14
1640 MW;
                                                                                                                                                                                 1 AGCKNFFWKTFTSC 14
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SEQUENCE OF 82-114 FROM N.A.
      [5]
SEQUENCE OF 79-92.
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             Primary structures of somatostatins from the islet organ of the haffish suggest an anomalous pathway of posttranslational processing processing processing 212:1855-1859(1988).

Endocrinology 122:1855-1859(1988).

I PUNCTION: SOMATOSTATIN INHIBITS THE RELEASE OF SOMATOTROPIN.

SUBCELLUTAR LOCATION: Socreted.

SIMILARITY: BELONGS TO THE SOMATOSTATIN FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Schally A.V., Huang W.-Y., Chang R.C.C., Arimura A., Redding T.W., Millar R.P., Hunkapiller M.W., Hood L.E., Isolation and structure of pro-somatostatin: a putative somatostatin precursor from ply hypothalamus." Proc. Natl. Acad. Sci. U.S.A. 77:4489-4493(1980).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Isolation and characterization of proSS1-32, a peptide derived from the N-terminal region of porcine preprosomatostatin.", EBS Lett. 192:141-146(1985).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             21-JUL-1986 (Rel. 01, Created)
01-OCT-1989 (Rel. 12, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
Somatostatin precursor (Contains: Somatostatin-28; Somatostatin-14)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Enkaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Cetartiodactyla; Suina; Suidae; Sus.
(MSBI_TaxID=9823;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  $EQUENCE OF 1-64.

MEDLINE-89278131; PubMed-2567292;

MEDLINE-89278131; PubMed-2567292;

MEDLINE-89278131; PubMed-2567292;

MEDLINE 1-64 Ba major product of somatostatin gene expression in pancreas and gut.";

J. Blol. Chem. 264:10633-10636(1989).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDLINE-86030691; PubMed-2865169;
Schmidt W.E., Mutt V., Kratzin H., Carlquist M., Conlon J.M.,
Creutzfeldt W.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             radayrol L., Joernvall H., Mutt V., Ribet A.;
N-terminally extended somatostatin: the primary structure of
                                                                                                                                                                                                                                                                                                                                      Score 77; DB 1; Length 34; Pred. No. 1e-05;
                                                                                                                                                                                                                                                                                                                                                                              1; Indels
                                                                                                                                                                                                                                                                                                 54FF213AAF424C75 CRC64;
Conlon J.M., Askensten U., Falkmer S., Thim L.;
                                                                                                                                                                                                                                          SOMATOSTATIN-34.
SOMATOSTATIN-14.
                                                                                                                                                                               Pfam; PF03002; Somatostatin; 1.
Cleavage on pair of basic residues; Hormone.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  92 AA.
                                                                                                                                                                                                                                                                                                                                                                            1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRT;
                                                                                                                                                             IPR004250; Somatostatin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                EDLINE-80113258; PubMed-7353633;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 IISSUE=Hypothalamus;
MEDLINE=81054799; PubMed=6107906;
                                                                                                                                                                                                                                                                                                                                    85.78;
                                                                                                                                                                                                                                                                             23 34
· 34 AA; 3963 MW;
                                                                                                                                                                                                                                                                                                                                                                                                            1 AGCKNFYWKGFTSC 14
                                                                                                                                                                                                                                                                                                                                                                                                                                     21 AGCKNFFWKTFTSC 34
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        EBS Lett. 109:55-58(1980).
                                                                                                                                                                                                                                                                                                                   Ouery Match
Best Local Similarity, 85.7%
....hes 12; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STANDARD;
                                                                                                                                              PIR; A32271; A3227
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             EQUENCE OF 65-92.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            scrofa (Pig).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Fragment).
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                                                                                                                                                                InterPro;
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P01168;
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STETTE COCCERTER S
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                    Schally A.V., Dupont A., Arimura A., Redding T.W., Nishi N.,
Libthbloum G.L., Schlesdinger D.H.,
Lisolation and structure of somatostatin from porcine hypothalami.",
Biochemistry 15:509-514(1976).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Actinopterygii; Neopterygii; Teleostei; Ostariophysi; Siluriformes
                                                                                                                                                                                                Riquet J.;
Submitted (SEP-1995) to the EMBL/GenBank/DDBJ databases.
-I- FUNCTION: SOMATOSTATIN INHIBITS THE RELEASE OF SOMATOTROPIN.
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POLITY:
1-10TL-1996 (Rel. 01, Created)
01-NUG-1996 (Rel. 15, Last sequence update)
16-0CT-2001 (Rel. 40, Last annotation update)
Somatostatin I precursor (Contains: Somatostatin-14 (SS-14)].
Ictalurus punctus (Channel caffish)
Ictalurus Williams (Caniata; Vertebrata; Euteleost
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 77; DB 1; Length .-. Pred. No. 2.6e-05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  10346 MW; 787CBE82CFBBAE76 CRC64;
                                                                                                                                                                                                                                                                                  -1- SUBCELLULAR LOCATION: Secreted.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOMATOSTATIN-14.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOMATOSTATIN-28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Pfam; PF03002; Somatostatin; 1.
Sleavage on pair of basic residues; Hormone.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1; Mismatches
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MEDLINE-85303576; PubMed-2863931;
Dixon J.E., Andrews P.C.;
"Somatostatins of the channel catfish.";
Adv. Exp. Med. Biol. 188:19-29(1985).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            somatostatin-14 messenger RNA.";
J. B1ol. Chem. 257:10372-10377(1982).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   InterPro; IPR004250; Somatostatin, 75fam; PF03002; Somatostatin,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRT;
PubMed-1252409;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         EMBL; U36385; AAB38485.1; -.
                                                                                                                                                               SEQUENCE OF 22-92 FROM N.A.
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es 12; Conserv
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                                      This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         "Molecular cloning of the cDNAs and distribution of the mRNAs encoding two somatostatin precursors in the African lungfish Protopterus
                                                                                                                                                                                                                                                                                                                                                                    leavage on pair of basic residues; Hormone; Signal; Multigene family. IGNAL 124 POTENTIAL.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 eavage on pair of basic residues; Hormone; Multigene family; Signal.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Protopterus annectens (African lungfish),
Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Dipnoi, Lepidosireniformes, Protopteridae, Protopterus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDLINE-99326690; PubMed-10398054;
Trabucchi M., Tostivint H., Lihrmann I., Jegou S., Vallarino M.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Comp. Neurol. 410:643-652(1999).
1- FUNCTION: SOMATOSTATIN INHIBITS THE RELEASE OF SOMATOTROPIN.
1- SUBCELLUIAR LOCATION: Secreted.
1- SIMILARITY: BELONGS TO THE SOMATOSTATIN FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        16-OCT-2001 (Rel. 40, Created)
16-OCT-2001 (Rel. 40, Last sequence update)
16-OCT-2001 (Rel. 40, Last unotation update)
Somatostatin I precursor (PSS1) (Contains: Somatostatin-27;
                                                                                                                                                                                                                                                                                                                                                                                                                                      POTENTIAL.
SOMATOSTATIN-26 (POTENTIAL).
SOMATOSTATIN-14.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SOMATOSTATIN-27 (POTENTIAL) SOMATOSTATIN-14.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      B5920015E2D272A4 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 77; DB 1; I
Pred. No. 3.2e-05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRT; 115 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    POTENTIAL
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InterPro; IPR004250; Somatostatin.
Pfam; PF03002; Somatostatin; 1.
                                                                                                                                                                                                                                                                         EMBL; U40754; AAD09359.1; -.
InterPro; IPR004250; Somatostatin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          12574 MW;
                                                                                                                                                                                                                                                                                                                                                   Somatostatin;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        101 AGCKNFFWKTFTSC 114
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 AGCKNFYWKGFTSC 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 86.5
Best Local Similarity 85.7
Matches 12; Conservative
                                                                                                                                                                                                                                                                                                                   nterPro; IPR004250;
fam; PF03002; Somato
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          103 1
114 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NCBI_TaxID=7888;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Somatostatin I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SMS1_PROAN Q9W7F0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      EPTIDE .
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DISULFID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 14
SMS1_PROAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             g
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation. the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@lab-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Eukaryota; Metazoa; Chordatá; Craniata; Vertebrata; Euteleostomi;
exfinopterygii; Neopterygii; Teleostei; Ostariophysi; Cypriniformes;
Cyprinidae; Carassius
NCBL_TaxID-7957;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Lin X.-W., Peter R.B.; Cloning and characterization of CDNAs encoding preprosomatostatin-I
                                                                                                                                                                                                              MEDLINE-81264223; Pubmed-6114953; Andrews P.C., Dixon J.E., Andrews P.C., Dixon J.E., "Isolation and structure of a peptide hormone predicted from a mRNA sequence. A second somatostatin from the catfish pancreas."; J. Biol. Chem. 256:8267-8270(1981).
-1 - FUNCTION: SOMATOSTATIN INHIBITS THE RELEASE OF SOMATOTROPIN.
-1 - SUBSELLUIAR LOCATION: Secreted.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 16-OCT-2001 (Rel. 40, Created)
16-OCT-2001 (Rel. 40, Last sequence update)
16-OCT-2001 (Rel. 40, Last amortation update)
Somatostatin IA precursor [Contains: Somatostatin-26; Somatostatin-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          om goldfish brain.";
(NOV-1995) to the EMBL/GenBank/DDBJ databases.
ON: SOMATOSTATIN INHIBITS THE RELEASE OF SOMATOTROPIN.
MEDLINE-82082515; Pubmed-6171821;
Taylor W.L., Collier K.J., Deschenes R.J., Weith H.L., Dixon J.
"Sequence analysis of a cDNA coding for a pancreatic precursor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 77; DB 1; Length 114;
Pred. No. 3.2e-05;
!; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        62 62 E -> Q (IN REF. 2).
114 AA; 12419 MW; FEE0F2C76F74D99F CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        pair of basic residues; Hormone; Signal;
                                                                                                                                                                                                                                                                                                                                                                                                                            -1- TISSUE SPECIFICITY: Pancreas.
-1- SIMILARITY: BELONGS TO THE SOMATOSTATIN FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 -i - SUBCELLUIAR LOCATION: Secreted.
                                                                                                                      roc. Natl. Acad. Sci. U.S.A. 78:6694-6698(1981).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PROBABLE.
SOMATOSTATIN-14.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRT; 114 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Somatostatin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           86.5%;
ilarity 85.7%;
Conservative 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Carassius auratus (Goldfish).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Somatostatin;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       EMBL; M25903; AAA49339.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       101 AGCKNFFWKTFTSC 114
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 AGCKNFYWKGFTSC 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STANDARD;
                                                                                                                                                                                         SEQUENCE OF 101-114.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           12; Conserv
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Pfam; PF03002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SSUE-Brain;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SMSA_CARAU
Q9YGH5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Submitted
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SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Cleavage
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Best Local 8
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Matches
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THE PRINCE OF SERVICE OF SERVICE

Length 114;

August 13, 2003, 14:51:19

Length 115;

BOCEF1E603FEAF09 CRC64;

SEQUENCE 115 AA; 12600 MW;

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Search completed: Augus
Job time : 6.38462 secs
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TH., Chartrel N., Conlon J.M.; lation of [Pro2,Metl3]somatostatin-14 and somatostatin-14 from the brain reveals the existence of a somatostatin gene family in a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             of basic residues; Hormone; Multigene family; Signal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ina ridibunda (Laughing frog) (Marsh frog).
karyota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
phibia; Batrachia; Anura; Neobatrachia; Ranoidea; Ranidae; Rana
                                                                                                                                                                                                                                                                                                                                                                  -OCT-2001 (Rel. 40, Created)
CCT-2001 (Rel. 40, Last sequence update)
-OCT-2001 (Rel. 40, Last annotation update)
matostatin 1 precursor (PSS1) [Contains: Somatostatin-14 (S-I)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Bucharles C., Vieau D., Coulouarn Y.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ys. Res. Commun. 188:477-482(1992).
SOMATOSTATIN INHIBITS THE RELEASE OF SOMATOTROPIN.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Cocurrence or two somatostatin variants in the frog brain: haracterization of the CDNAs, distribution of the mRNAs, and eceptor binding affinities of the peptides."; roc. Natl. Acad. Sci. U.S.A. 93:12605-12610(1996).
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BY SIMILARITY.
349756FEB4ABE213 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         - SUBCELLULAR LOCATION: Secreted.
- SIMILARITY: BELONGS TO THE SOMATOSTATIN FAMILY.
   Score 77; DB 1; Pred. No. 3.2e-05;
                                                                                                                                                                                                                                                                                                                 115 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         EDLINE-93038702; PubMed-1358069;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  12691 MW;
Query Match
Best Local Similarity 85.7%;
Matches 12; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 86.5%;
Best Local Similarity 85.7%;
Matches 12; Conservative
                                                                                                                                                  102 AGCKNFFWKTFTSC 115
                                                                                                              1 AGCKNFYWKGFTSC 14
                                                                                                                                                                                                                                                                                                                 STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     EQUENCE OF 102-115.
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SMS1_RANRI
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#8888 #8888

102 AGCKNFFWKTFTSC 115

1 AGCKNFYWKGFTSC 14

à g Q8fee2 escherichia Q8foe3 drosophila Q8h7a7 arabidopsis Q8h7a7 arabidopsis Q8h015 cryza sativ Q9culc m adult mal Q6592 aorine ant Q6592 aprine art Q92ba3 neisseria m Q9fav6 drosophila Q60958 leishemnia Q9far1 arabidopsis Q9fitla arabidopsis Q19698 ceenorhabdi

201

Run on:

28 AA.

ALIGNMENTS

Gaps Carassius auratus (Goldfish).
Eukaryota: Metazoa: Chordata: Craniata: Vertebrata: Euteleostomi;
Actinopterygil: Neopterygil: Teleostei; Ostariophysi; Cypriniformes;
Cyprinidae; Carassius.
NCBL_TaxID=7957; Uesaka T., Yano K., Yamasaki M., Ando M.; "Somatostatin-, vasoactive intestinal peptide-, and granulin-like peptides isolated from intestinal extracts of goldfish, Carassius ő Score 89; DB 13; Length 28; Pred. No. 6.2e-08; Mismatches 0; Indels 15D271F677C945BE CRC64; Created) Last sequence update) Last annotation update) Comp. Endocrinol. 99:298-306(1995). Pro; IPR004250; Somatostatin. 090141 PRELIMINARY; PRT; 090141; 01-DEC-2001 (TEMBLEL 19, Created) GSS-28-SOMATOSTATIN-11ke peptide. Carassius auratus (Goldfish). MEDLINE-96051491; PubMed-8536941; ö Pfam; PF03002; Somatostatin; 1 SEQUENCE 28 AA; 3204 MW; 1 Ouery Match 100.0%; Best Local Similarity 100.0%; Matches 14; Conservative 0; RESULT 2. Q90Y41 ID Q90Y4 AC Q90Y4 DT 01-DE ò

Q8luv6 homo saplen Q8nfe5 homo saplen Q9prv6 anguilla ja Q42113 brachydanio Q8xf27 salmonella

brachydanio 090y39 catostomus 090xe0 acipenser t

09DDE4

Q90y41 gnathonemus Q90y40 chitala chi Q90y42 pantodon bu Q90y43 osteoglossu

Result

Ogprvo anguilla ja Ogjhx5 brachydanio 090xel acipenser t

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01-MAY-2000 (TrEMBLrel. 13, Created)
01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-DBC-2001 (TrEMBLrel. 19, Last annotation update)
somatostatin-related peptide.
somatostatin-related peptide.
by appuila japonica (Japanese eel).
Eukaryota: Metazoa: Chordata: Craniata; Vertebrata: Buteleostomi;
Actinopterygii; Neopterygii: Teleostei; Anguilliformes; Anguillidae.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ol-Mar. vova ...
Preprosomatostatin.
Osteoglossum bicirrhosum (silver arawana).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Eukaryota; Metazoa; Chordata; Teleostei; Osteoglossomorpha;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ó
                                                                                                                                                   Al-Mahrouki A.A., Irwin D.M., Youson J.H.; Characterization of variant somatostatin cDNAs from several osteoglossomorphs: molecular identification and comparative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           "Characterization of variant somatostatin cDNAs from several osteoglossomorphs: molecular identification and comparative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 115;
   Actinopterygii; Neopterygii; Teleostei; Osteoglossomorpha; Osteoglossiformes; Pantodontidae; Pantodon. NCBL_TaxID-8276;
                                                                                                                                                                                                                                                                           Submitted (AUG-2000) to the EMBL/GenBank/DDBJ databases.
EMBL; AF292651; AAK97068.1; -
InterPro: JPR004250; Somartostatin.
Pfan; PPT0925, Somartostatin; 1.
SEQUENCE 114 AA; 12352 MW; 7E3D44CB6A27B12F CRC64;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Submitted (AUG-2000) to the EMBL/GenBank/DDBJ databases.
EMBL; AF292650; AAK97067.1; -
Interpro; IPR004250; Somatostatin.
Pran; PR0302; Somatostatin; 1.
SEQUENCE 115 AA; 12791 MW; D65FBD7C6F1E4E4D CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       01-DEC-2001 (TrEMBLrel. 19, Created)
01-DEC-2001 (TrEMBLrel. 19, Last sequence update)
01-MAR-2002 (TrEMBLrel. 20, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 89; DB 13;
Pred. No. 2.5e-07;
0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 89; DB 13;
Pred. No. 2.5e-07
; Mismatches 0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Al-Mahrouki A.A., Irwin D.M., Youson J.H.; "Characterization of variant somatostatin of standard somatostatin of standard sometostatin of standard sometostatin s
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRT;
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Local Similarity 100.0%;
les 14; Conservative C
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Ouery Match
Best Local Similarity 100.0%;
Matches 14; Conservative (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              102 AGCKNFYWKGFTSC 115
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE FROM N.A.
                                                                                                                              SEQUENCE FROM N.A.
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NCBI_TaxID=7937;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 6
Q9PRV0
          δ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0; Gaps
                                                                                                                       Bukaryota; Metazoa; Chordata; Cranlata; Vertebrata; Euteleostomi;
Actinopteryqli, Neopteryglij Teleostei; Osteoglossomorpha;
Gsteoglossiformes; Mormyrldae; Gnathonemus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               chitala chitala (clown knifefish).
Bukaryota; Metazoa; Choródata; Craniata; Vertebrata; Euteleostomi; detinopteryota; Neopterygii; Teleostei; Osteoglossomorpha; dsteoglossiformes; Notopteridae; Chitala.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 89; DB 13; Length 114;
Pred. No. 2.5e-07;
; Mismatches 0; Indels 0
                                                                                                                                                                                                                                                                                                    Al Mahrouki A.A., Irwin D.M., Youson J.H.; Alcharacterization of Variant somatostatin cDNAs from several esteoglossomorphs: molecular identification and comparative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    . Match 100.0%; Score 89; DB 13; Length 114; Local Similarity 100.0%; Pred. No. 2.5e-07; es 14; Conservative 0; Mismatches 0; Indels (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Al-Mahrouki A.A., Irwin D.M., Youson J.H.,
Characterization of variant somatostatin conna erveral
deteoglossomorphs: molecular identification and comparative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Submitted (AUG-2000) to the EMBL/GenBank/DDBJ databases.
EMBL, AF792653; AAK97070.1; -
InterPro; IPR004250; Somatostatin.
Ffam; PF03002; Somatostatin; 1.
SEQUENCE 114 AA; 12561 MW; 4E3C32F58E34F971 CRC64;
                                                                                                                                                                                                                                                                                                                            dentification and compara communitied (AUG-2000) to the EMBL/GenBank/DDBJ databases interpro: IPR004250; Somatostatin. Ffam; PP03002; Somatostatin. SEQUENCE 114 a.m.
-DEC-2001 (TrEMBLrel. 19, Last sequence update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    -DEC-2001 (TrEMBLrel. 19, Created)
-DEC-2001 (TrEMBLrel. 19, Last sequence update)
-JUN-2002 (TrEMBLrel. 21, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Created)
Last sequence update)
Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRT; 114 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Pantodon buchholtzi (Butterflyfish)
Bukaryota; Metazoa; Chordata; Crani
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      101 AGCKNFYWKGFTSC 114
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRELIMINARY;
                                                                                               athonemus petersil.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Preprosomatostatin.
Pantodon buchholtzi
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                                                                  reprosomatostatin
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                                                                                                                                                                                                                                                                           EQUENCE FROM N.A.
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                                                                                                                                                                                                               NCBI_TaxID-42645;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query, Match
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090x40;
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transmontanus; molecular cloning and distribution of the mRNAs encoding two somatostatin precursors.";
- Comp. Neurol. 0:0-0(2001).
                                                                                                                                                   AAL13248.1; -. 4250; Somatostatin.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Dores R.M., Vaudry H.,
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                                                                                                             J. Comp. Neurol
EMBL, AF395849;
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                                 Desaka T., Yano, K., Yamasaki M., Nagashima K., Ando M.; Somatostatin.related peptides isolated from the eel gut: effects on ion and water absorption across the intestine of the seawater eel.", T. Exp. Biol. 188:205-216(1994).

InterPro: IPR004250; Somatostatin.

PFGam; PF03002; Somatostatin; BFC672143A04A3F5 CRC64;
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Brachydanio rerio (Zebrafish) (Danio rerio).
Brkaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Actinopterygii; Neopterygii; Teleostei; Ostariophysi; Cypriniformës;
NCBI_TaxID-7955;
                                                                                                                                                                                                                                                                                                                                                                                                                                                          0; Gaps
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Mech. Dev. 115:133-137(2002).

Inter Pro. 1 PR004250; Somatostatin.

Pfam; PF03002; Somatostatin.

Pfam; PF03002; Somatostatin.

SEQUENCE 114 AA; 12416 MW; 3D41424AE54E74C8 CRC64;
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Actinopterygil; Chondrostel; Acipenseriformes; Acipenseridae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Tostivint H., Lihrmann I., Sollars C., Vallarino M.
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Prabucchi M. Tostivint H., Lihrmann I., Sollars C., Vallarino Mores R.M., Vaudry H.;
Polygenic expression of somatostatin in the sturgeon Acipenser
                                                                                                                                                                                                                                                                                                                                                           Query Match 88.8%; Score 79; DB 13; Length 25; Best Local Similarity 92.9%; Pred. No. 2.6e-06; Matches 13; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         01-0cr-2002 (TrEMBLrel. 22, Created)
01-0cr-2002 (TrEMBLrel. 22, Last sequence update)
01-MAR-2003 (TrEMBLrel. 23, Last annotation update)
Somatostatin-14.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              01-DEC-2001 (TrEMBLrel. 19, Created)
01-DEC-2001 (TrEMBLrel. 19, Last sequence update)
01-MAR-2002 (TrEMBLrel. 20, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRT; 114 AA.
MEDLINE-95053622; PubMed-7525832;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRELIMINARY;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Acipenser
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Q8JHX5
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090 x21
10 090 x21
10 090 x21
10 010 x21
10 010 x21
10 x
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01-DEC-2001 (TTEMBLrel. 19, Created)
01-DEC-2001 (TTEMBLrel. 19, Last sequence update)
01-MAR-2002 (TTEMBLrel. 20, Last annotation update)
01-MAR-2002 (TTEMBLRel. 20, Last annotation update)
Catcostomus commercent (White sucker)
Ekkaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Catcostomidae; Catostomus
Catcostomidae; Catostomus
Catcostomidae; Catostomus
Catcostomidae; Catostomus
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Trabucchi M., Tostivint H., Lihrmann I., Sollars C., Vallarino M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            "Polygenic expression of somatostatin in the sturgeon Acipenser transmontanus: molecular cloning and distribution of the mRNAs encoding two somatostatin precursors.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           090XE0;
01-DEC-2001 (TrEMBLrel. 19, Last sequence update)
01-DEC-2001 (TrEMBLrel. 20, Last sequence update)
01-MAR-2002 (TrEMBLrel. 20, Last annotation update)
Somatostatin pro2.
Acipense transmontanus (White sturgeon).
Bukaryota, Metazoa; Chordata, Craniata; Vertebrata; Euteleost
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match

86.58; Score 77; DB 13; Length 120;
Best Local Similarity 85.78; Pred No. 2.6e-05;
Matches 12; Conservative 1; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE FROM N.A.
Al-Mahrouki A.A., Irwin D.M., Youson J.H.;
"Molecular cloning and characterization of white sucker preprosomatostatin.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Submitted (AUG-2000) to the EMBL/GenBank/DDBJ databases. EMBL: AF292654; AAK97071.1; Interpro; IPR004250; Somatostatin. Pfam: PF03002; Somatostatin; 1. SEQUENCE 120 AA; 13783 MW; 00828D35263E8805 CRC64;
InterPro; IPR004250; Somatostatin.
Pfam; PF03002; Somatostatin; 1.
SEQUENCE 116 AA; 12616 MW; 72E0C3FF6C80650F CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRT; 120 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRT; 111 AA.
                                                                                                                                                                                                                                                                                     103 AGCKNFFWKTFTSC 116
                                                                                                                    Query Match
Best Local Similarity 85.7%;
Matches 12; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             107 AGCKNFFWKTFTSC 120
                                                                                                                                                                                                                                                   1 AGCKNFYWKGFTSC 14
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Ol-MAY-2000 (TrEMBLrel. 13, Created)
01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-MAY-2002 (TrEMBLrel. 21, Last annotation update)
Somatostatin homolog (Fregment)
Somatostatin homolog (Fregment)
Somatostatin homolog (Treament)
Somatostatin homolog (Creament)
Somatostatin homolog (Creament)
Somatostatin homolog (Treament)
S
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OBNEES; O. (TERBILE). 22, Created)
01-0CT-2002 (TERBILE). 22, Last sequence update)
01-0CT-2003 (TERBILE). 23, Last annotation update)
Hypothetical protein.
Hypothetical protein.
Hypothetical protein.
Home sapiens (Human).
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Best Local Similarity 90.0%; Pred. No. 0.11;
Matches 9; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Cal Or Guo J. H. Yu L.,
Submitted (Jun-2002) to the EMBL/GenBank/DDBJ databases.
Submitted (Jun-2002) to the EMBL/GenBank/DDBJ databases.
EMBL., AF25016; AAM70482.1;
InterPro; IPR004822, Hattone_Core.
InterPro; IPR004250; Somatostatin.
Ffan; PF03002; Somatostatin; 1.
Hypothetical protein Protein Protein SEQUENCE 164 AA; 19047 MW; 389682C41252426A CRC64;
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SEQUENCE 23 AA; 2655 MW; BA4317DFF3BDBD29 CRC64;
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MEDLINE-95003944; PubMed-7765422;
UGSaka, I., Yano K., Yamasaki M., Ando M.;
Zool. Sci. 11.491-494(1994).
NON_TER
Matches 8; Conservative 3; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           152 CRNFFWKTFSSC 163
                                                                                                                                                                                              110 CRNFFWKTFSSC 121
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRELIMINARY;
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NCBI_TaxID-7937;
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O9PRV(
ID 09PRV(
ID 09PRV(
DT 01-MA
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QBNFE5
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01-Mar-2003 (TrEMBLrel. 23, Last sequence update)
01-Mar-2003 (TrEMBLrel. 23, Last annotation update)
51milar to cortistatin (Fragment).
Homo sapiens (Human)
BURATOGE, Merazoa, Chordata, Craniata, Vertebrata, Buteleostomi,
Mammalia: Butheria, Primates; Catarrhini, Hominidae, Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TISSUE-Pancreas;
MEDITE-99452919.
MEDITE-99452919.
MEDITE-9945190: PubMed-10495291;
**Argenton F., Zecchin E., Bortchussi M.;
**Early appearance of pancreatic hormone-expressing cells in the zebrafish embryo.**;
**Mech. Dev. 87:217-221(1999)
**EMBL; AJ238017; CAC20110.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Somatostatin.

SMST OR SOM.

Brachydanio rerio (Zebrafish) (Danio rerio).

Braryota, Metazoa, Chordata; Craniata; Vertebrata; Euteleostomi; Actinopteryqli; Neopteryqli; Teleostei; Ostariophysi; Cypriniform (Cyprinidae; Danio NCBL_TAXID=7955;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity 61.5%; Pred. No. 0.016;
Matches 8; Conservative 3; Mismatches 2; Indels
                                                                                                                                                                                        Query Match
Best Local Similarity 78.6%; Pred. No. 0.00053;
Matches 11; Conservative 1; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Strausberg R.; Submitted (NOV-2002) to the EMBL/GenBank/DDBJ databases.
EMBL; BOG40034.1; -1 NON_TER 1 1 SEQUENCE 122 AA; 13369 MW; A1279CA09CB0CB44 CRC64;
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EMBL, AJ338017. CACCHIO.1.
EMBL, AJ338017. CACCHIO.1.
InterPro, IPR004256; Somatostatin.
Pfam, PP03002; Somatostatin.
PGRORECE 107 AA; 11839 MW; E12C923E56642EFB CRC64;
                       InterPro; IPR004250; Somatostatin.
Pfam, PF03002; Somatostatin; 1.
SEQUENCE 111 A4, 12748 NW; 4E27DB90896A9025 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     01-WAR-2001 (TrEMBLrel. 16, Created)
01-WAR-2001 (TREMBLrel. 16, Last sequence update)
01-OCT-2002 (TREMBLrel. 22, Last annotation update),
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98 APCKNFFWKTFTSC 111
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Best Local Similarity
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ID Q9DDE4
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Search completed: August 13, 2003, 14:52:58 Job time : 22.1795 secs

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